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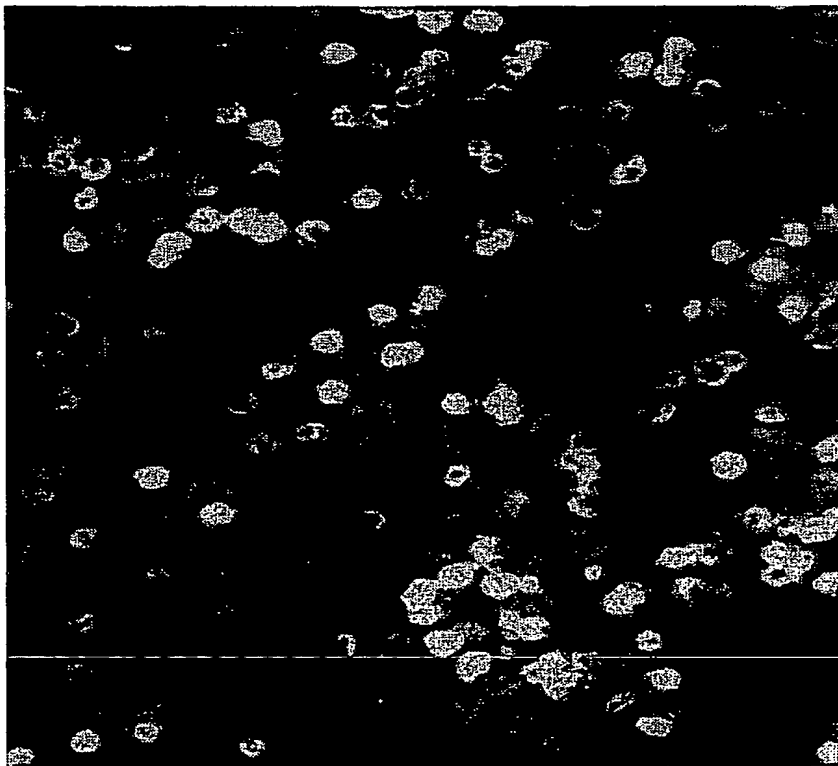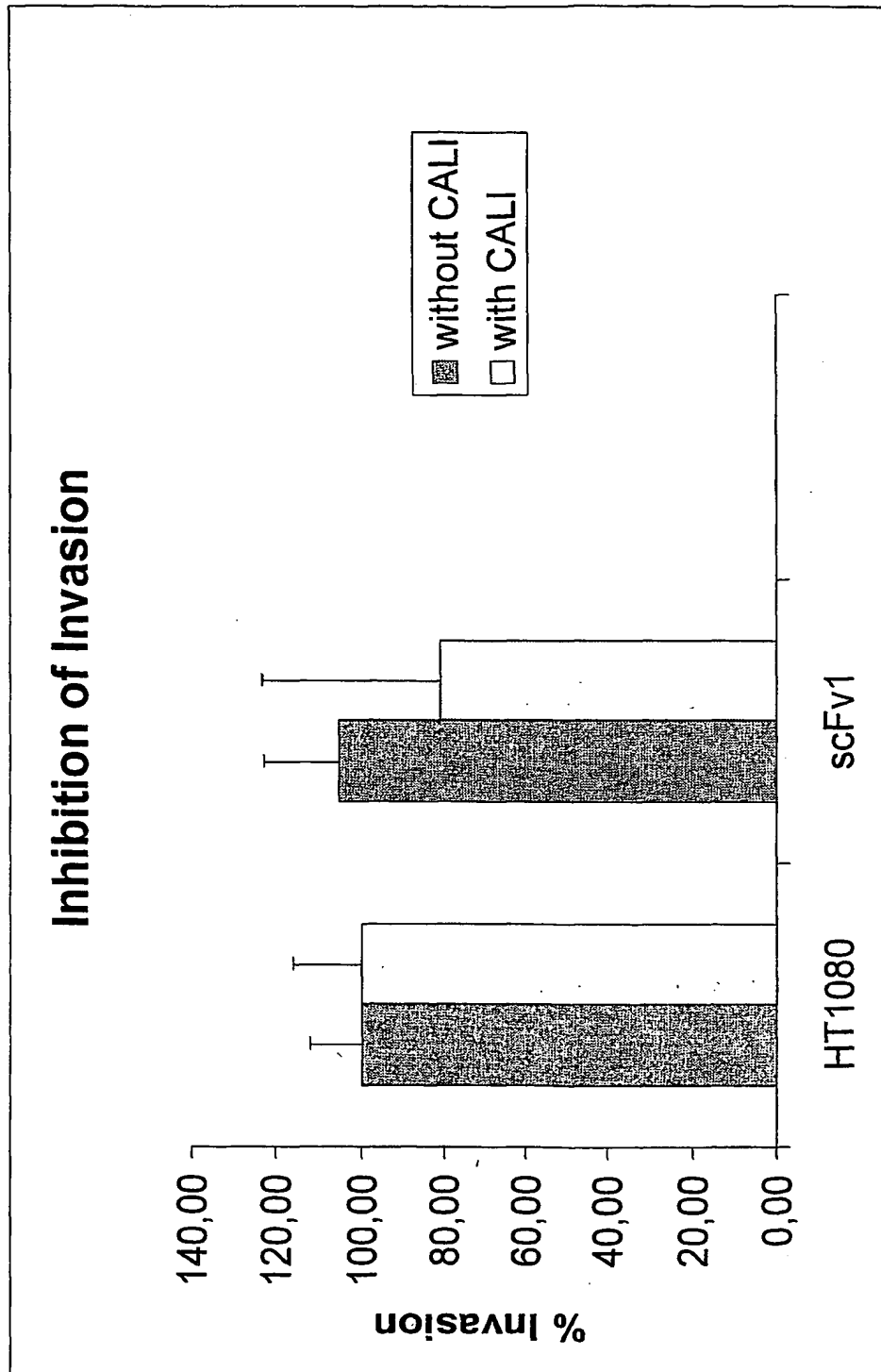

Invasion of Hs-27 control cells

Invasion of HT1080 cells
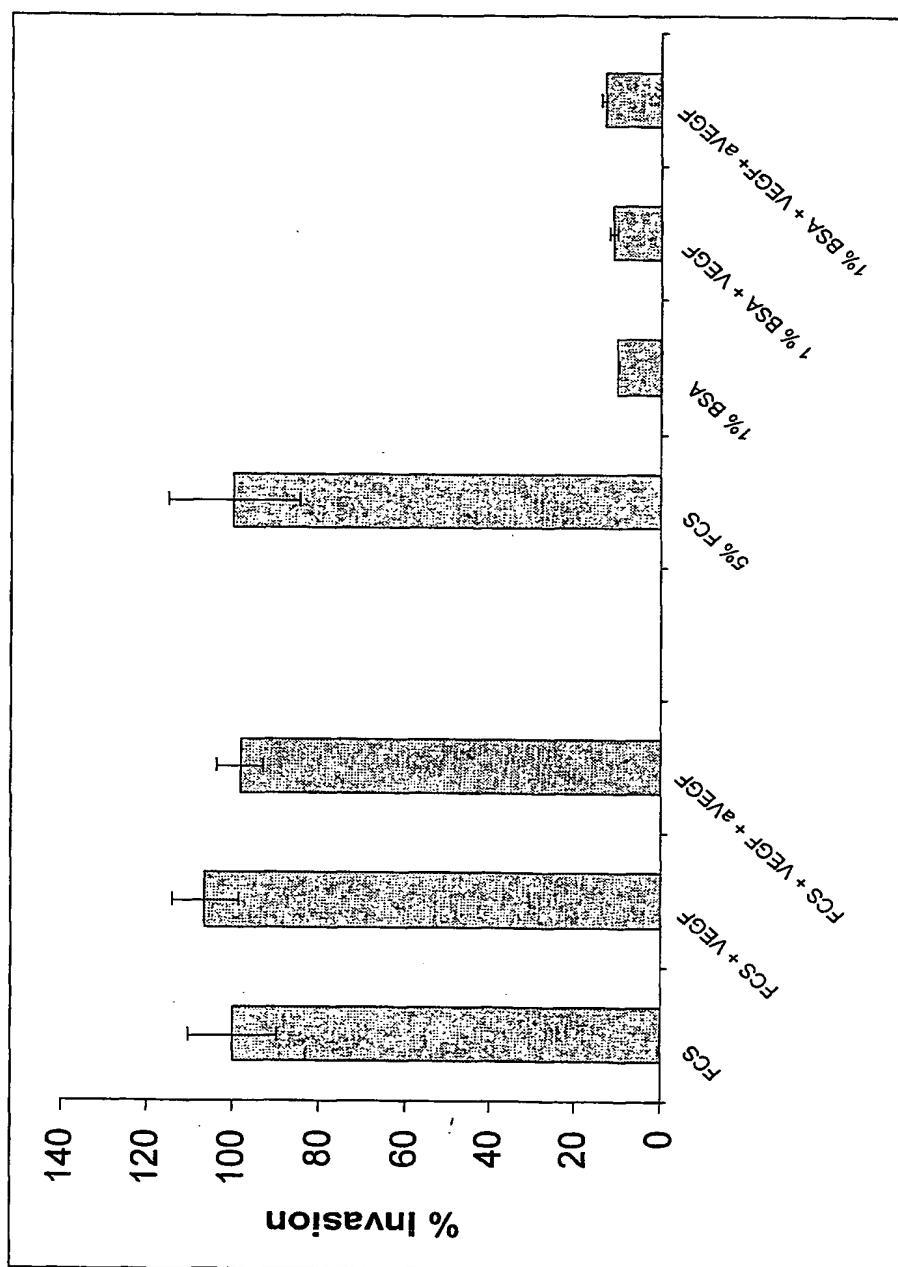
Fig. 1
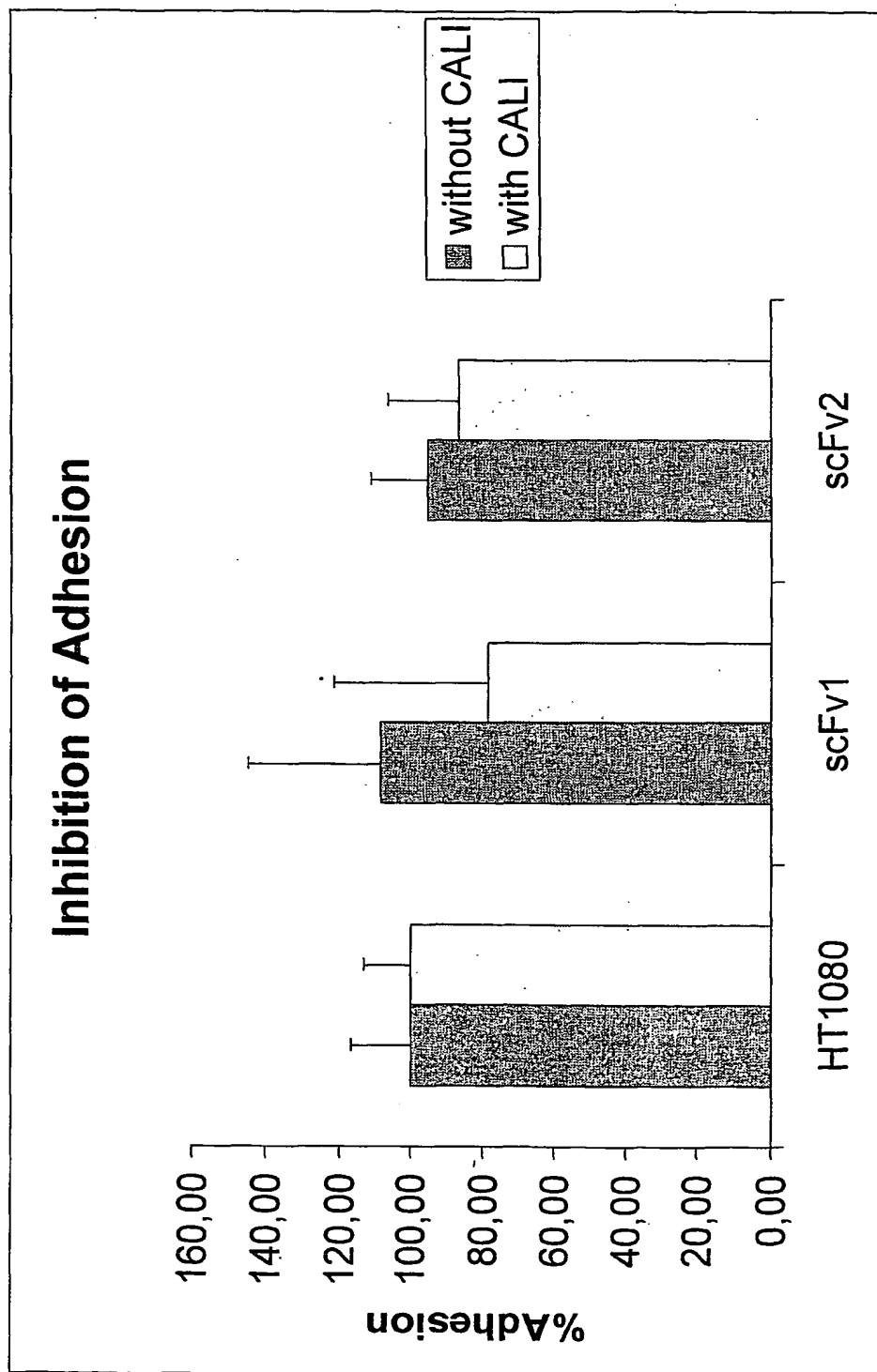
Fig. 2

Fig. 2a

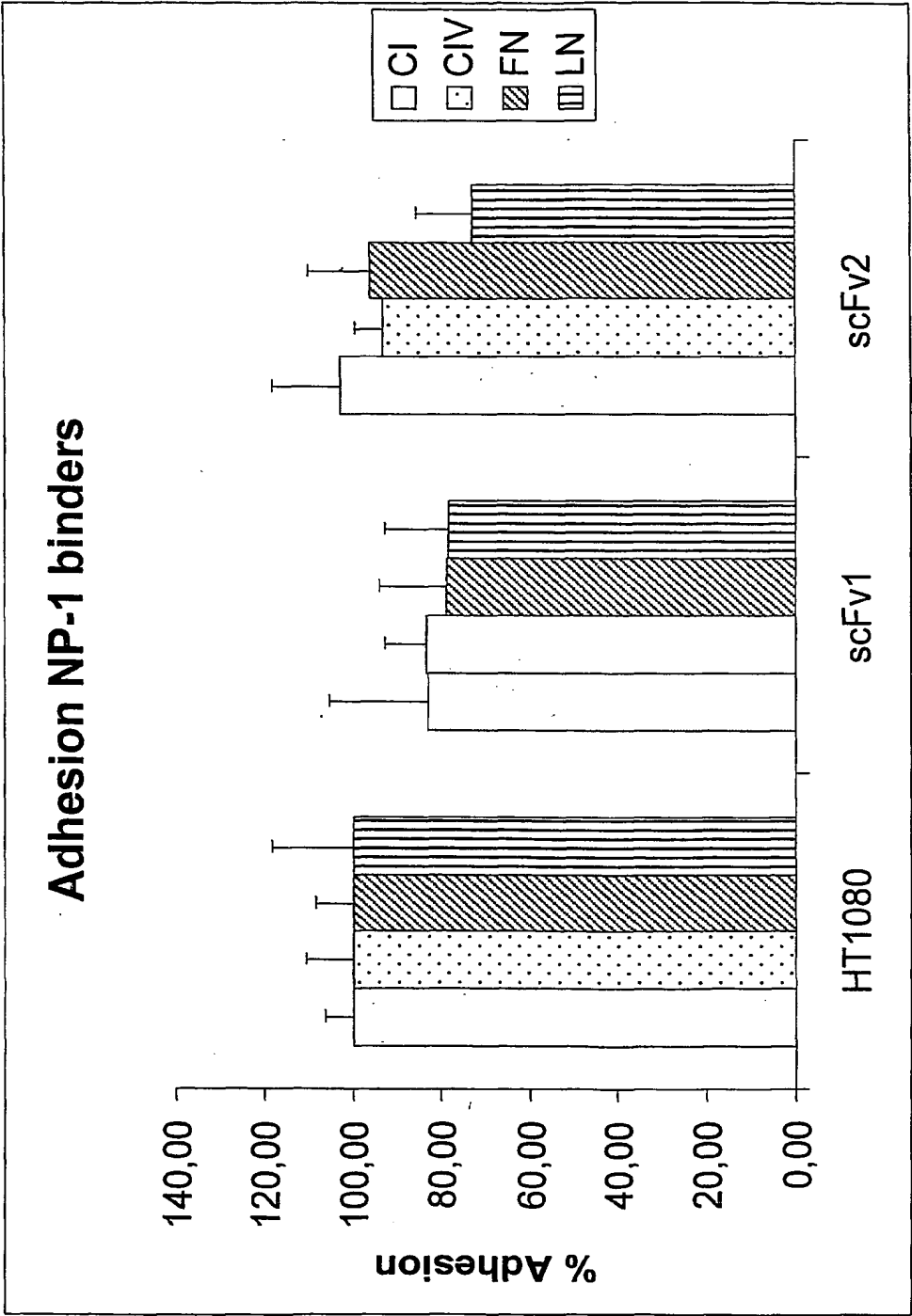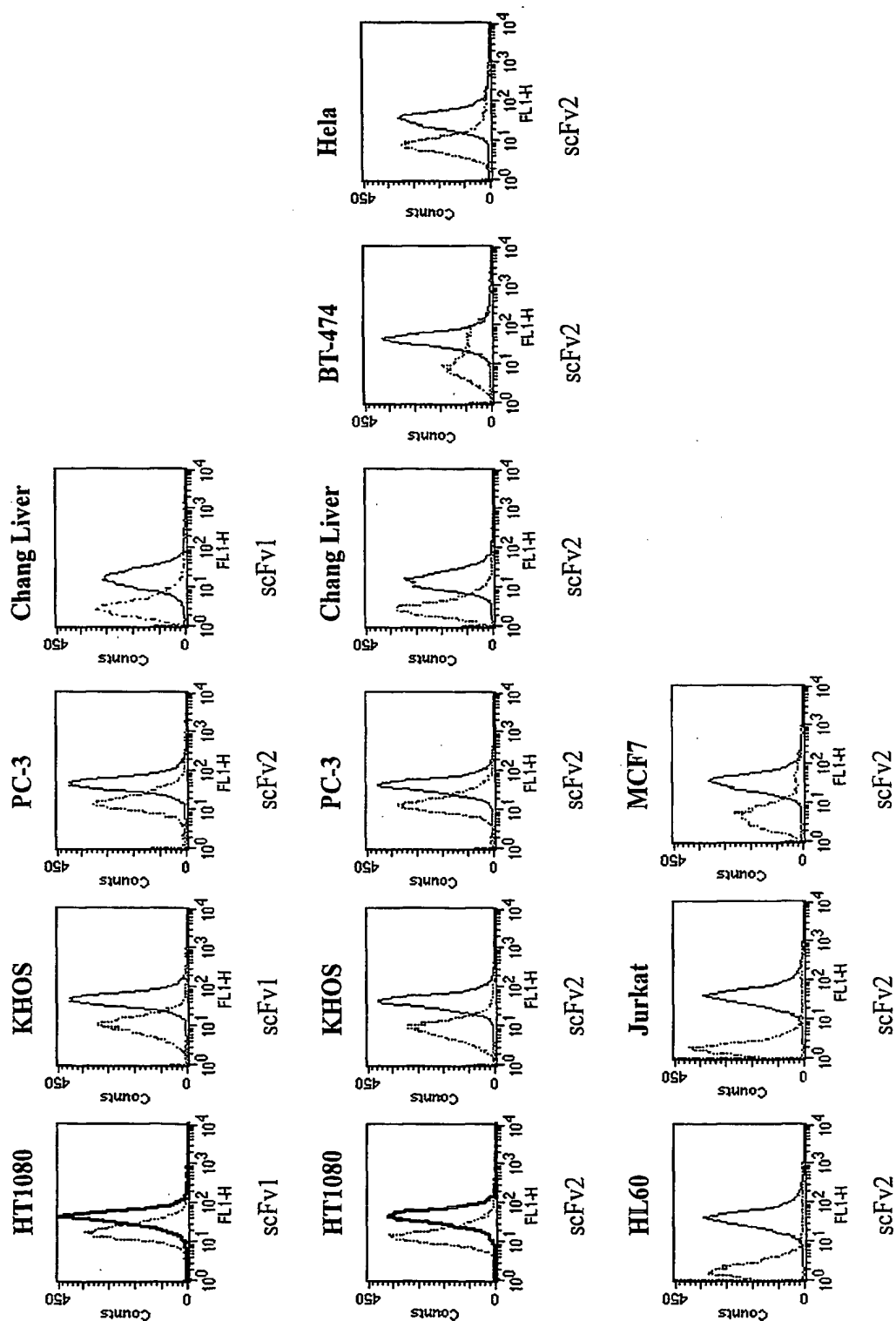Fig. 3

Fig. 4

Fig. 5

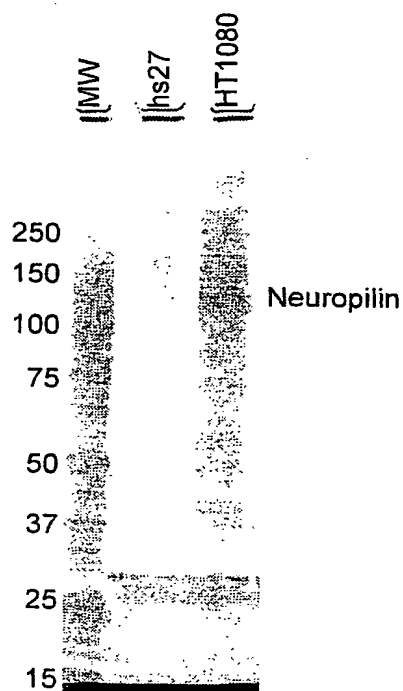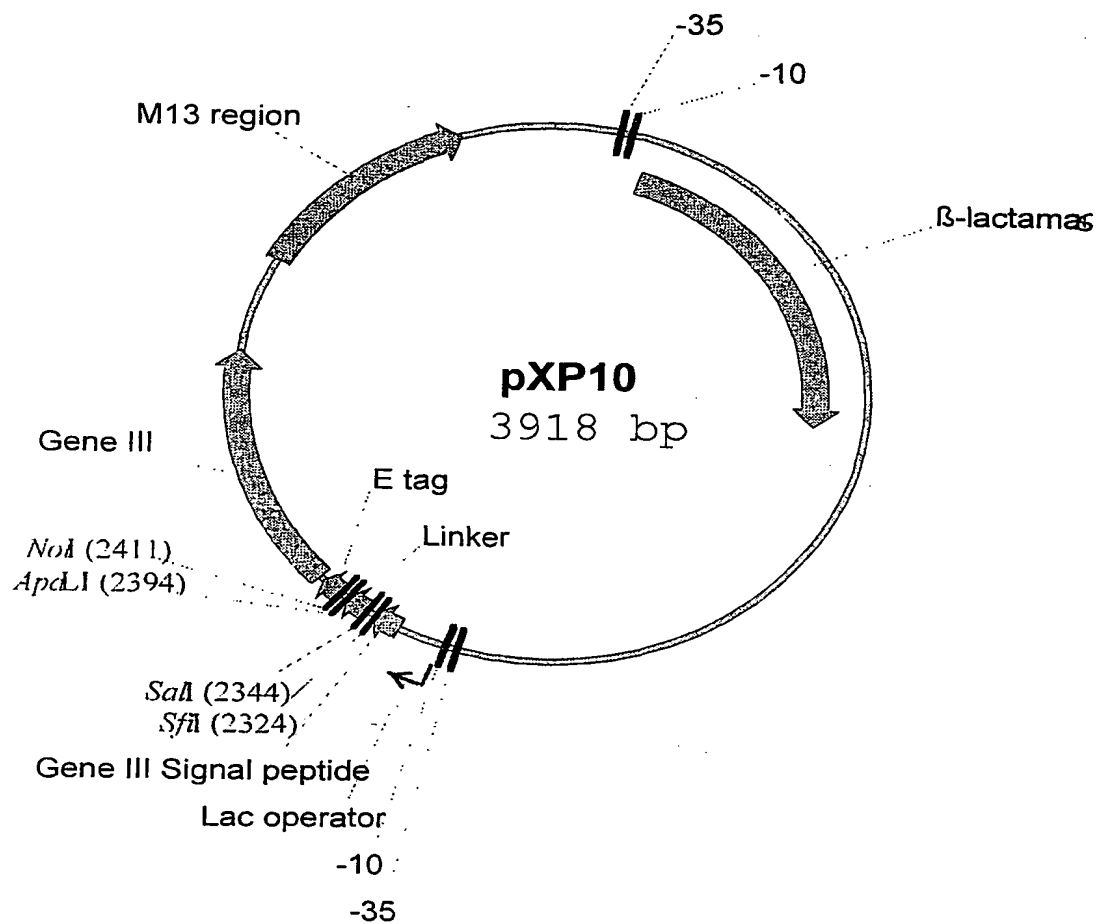# ~ 130 kDa = Neuropilin

Fig. 6

Fig. 7a

Nucleotide Sequences pXP10

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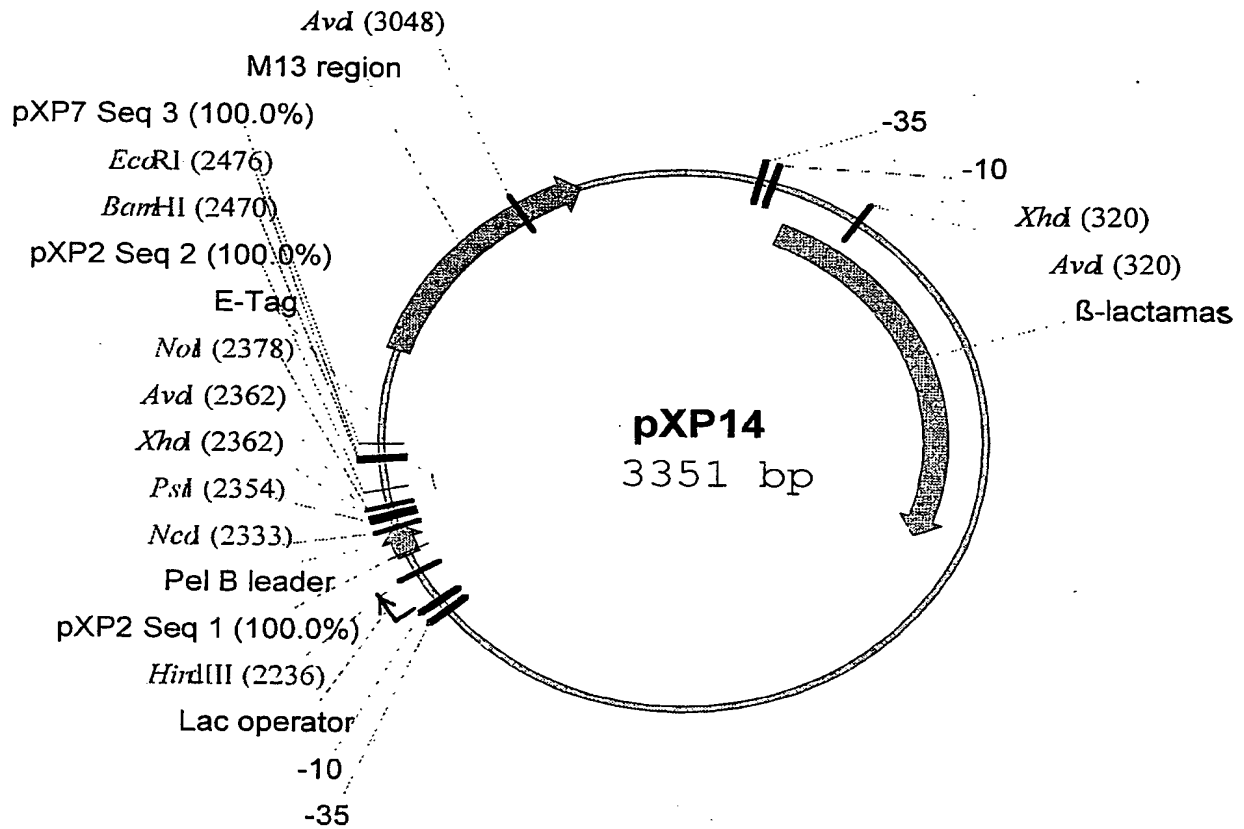
Fig. 7b

Fig. 8a

Nucleotide Sequences pXP14

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SUBSTITUTE SHEET (RULE 26)

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2201  TCACACAGGA  AACAGCTATG  ACCATGATTA  CGCCAAGCTT  GCATGCAAAT
      AGTGTGTCCT  TTGTCGATAC  TGGTACTAAT  GCGGTTCGAA  CGTACGTTTA
2251  TCTATTTCAA  GGAGACAGTC  ATAATGAAAT  ACCTATTGCC  TACGGCAGCC
      AGATAAAGTT  CCTCTGTCAG  TATTACTTTA  TGGATAACGG  ATGCCGTCGG
2301  GCTGGATTGT  TATTACTCGC  GGCCCAGCCG  GCCATGGCCC  AGGTGCAGCT
      CGACCTAACA  ATAATGAGCG  CCGGGTCGGC  CGGTACCGGG  TCCACGTCGA
2351  GCAGGTCGGC  CTCGAGATCA  AACGGGCGGC  CGCAGGTGCG  CCGGTGCCGT
      CGTCCAGCCG  GAGCTCTAGT  TTGCCCGCCG  GCGTCCACGC  GGCCACGGCA
2401  ATCCAGATCC  GCTGGAACCG  CGTGGGGCCG  CAAGCGCTTG  GAGCCACCCG
      TAGGTCTAGG  CGACCTTGGC  GCACCCCGGC  GTTCGCGAAC  CTCGGTGGGC
2451  CAGTTCGAAA  AATAATAAGG  ATCCGAATTC  ACTGGCCGTC  GTTTTACAAC
      GTCAAGCTTT  TTATTATTCC  TAGGCTTAAG  TGACCGGCAG  CAAAATGTTG
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2501   GTCGTGACTG   GGAAAACCCT   GGCGTTACCC   AACTTAATCG   CCTTGCAGCA
       CAGCACTGAC   CCTTTTGGGA   CCGCAATGGG   TTGAATTAGC   GGAACGTCGT
2551   CATCCCCCTT   TCGCCAGCTG   GCGTAATAGC   GAAGAGGCCC   GCACCGATCG
       GTAGGGGGAA   AGCGGTCGAC   CGCATTATCG   CTTCTCCGGG   CGTGGCTAGC
2601   CCCTTCCCAA   CAGTTGCGCA   GCCTGAATGG   CGAATGGCGC   CTGATGCGGT
       GGGAAGGGTT   GTCAACGCGT   CGGACTTACC   GCTTACCGCG   GACTACGCCA
2651   ATTTTCTCCT   TACGCATCTG   TGCGGTATTT   CACACCGCAT   ACGTCAAAGC
       TAAAAGAGGA   ATGCGTAGAC   ACGCCATAAA   GTGTGGCGTA   TGCAGTTTCG
2701   AACCATAGTA   CGCGCCCTGT   AGCGGCGCAT   TAAGCCCGGC   GGGTGTGGTG
       TTGGTATCAT   GCGCGGGACA   TCGCCGCGTA   ATTCGGGCCG   CCCACACCAC
2751   GTTACGCGCA   GCGTGACCGC   TACACTTGCC   AGCGCCCTAG   CCCCCGCTCC
       CAATGCGCGT   CGCACTGGCG   ATGTGAACGG   TCGCGGGATC   GGGGGCGAGG
2801   TTTCGCTTTC   TTCCCTTCCT   TTCTCGCCAC   GTTCGCCGGC   TTTCCCCGTC
       AAAGCGAAAG   AAGGGAAGGA   AAGAGCGGTG   CAAGCGGCCG   AAAGGGGCAG
2851   AAGCTCTAAA   TCGGGGGCTC   CCTTTAGGGT   TCCGATTTAG   TGCTTTACGG
       TTCGAGATTT   AGCCCCCGAG   GGAAATCCCA   AGGCTAAATC   ACGAAATGCC
2901   CACCTCGACC   CCAAAAAACT   TGATTTGGGT   GATGGTTCAC   GTAGTGGGCC
       GTGGAGCTGG   GGTTTTTTGA   ACTAAACCCA   CTACCAAGTG   CATCACCCGG
2951   ATCGCCCTGA   TAGACGGTTT   TTCGTCCTTT   GACGTTCGAG   TCCACGTTCT
       TAGCGGGACT   ATCTGCCAAA   AAGCAGGAAA   CTGCAAGCTC   AGGTGCAAGA
3001   TTAATAGTGG   ACTCTTGTTC   CAAACTGGAA   CAATACTCAA   CCCTATCTCG
       AATTATCACC   TGAGAACAAG   GTTTGACCTT   GTTATGAGTT   GGGATAGAGC
3051   GGCTATTCTT   TTGATTTATA   AGGGATTTTG   CCGATTTCGG   CCTATTGGTT
       CCGATAAGAA   AACTAAATAT   TCCCTAAAAC   GGCTAAAGCC   GGATAACCAA
3101   AAAAAATGAG   CTGATTTAAC   AAAAATTTAA   CGCGAATTTT   AACAAAATAT
       TTTTTTACTC   GACTAAATTG   TTTTTAAATT   GCGCTTAAAA   TTGTTTTATA
3151   TAACGTTTAC   AATTTTATGG   TGCAGTCTCA   GTACAATCTG   CTCTGATGCC
       ATTGCAAATG   TTAAATACC   ACGTCAGAGT   CATGTTAGAC   GAGACTACGG
3201   GCATAGTTAA   GCCAGCCCCG   ACACCCGCCA   ACACCCGCTG   ACGCGCCCTG
       CGTATCAATT   CGGTCGGGGC   TGTGGGCGGT   TGTGGGCGAC   TGCGCGGGAC
3251   ACGGGCTTGT   CTGCTCCCGG   CATCCGCTTA   CAGACAAGCT   GTGACCGTCT
       TGCCCGAACA   GACGAGGGCC   GTAGGCGAAT   GTCTGTTCGA   CACTGGCAGA
3301   CCGGGAGCTG   CATGTGTCAG   AGGTTTTCAC   CGTCATCACC   GAAACGCGCG
       GGCCCTCGAC   GTACACAGTC   TCCAAAAGTG   GCAGTAGTGG   CTTTGCGCGC
3351   A
       T
```

Fig. 8b

cDNA primers

VLK-c	CTGGATGGTGGGAAGATGGA
VLL-c	TCAGAGGAAGGAAACAGGGT
IgG1-c	CTTACAACCACAATCCCTGGGCACAATTTT
IgG2a-c	CTTTGTGGGCCCTCTGGGCTCAAT
IgG2b	TGAAATGGGCCCGCTGGGCTCAAG
IgG3-c	GGGCTTGGGTATTCTAGGCTCGAT

VH forward primers without restriction sites

M-VH1	GAGGTGCAGCTTCAGGAGTCAGG
M-VH2	CAGGTGCAGCTGAAGGAGTCAGG
M-VH3	GAGGTCCAGCTGCAACAGTCTGG
M-VH4	GAGGTTCAGCTGCAGCAGTCTGG
M-VH5	CAGGTCCAACTGCAGCAGCCTGG
M-VH6	CAGGTTCAGCTGCAGCAGTCTGG
M-VH7	GAGGTGAAGCTGGTGGAGTCTGG
M-VH8	GAGGTGAAGCTGGTGGAATCTGG
M-VH9	GAGGTTCAGCTTCAGCAGTCTGG

VH backward primers without restriction sites

M-JH1	TGAGGAGACGGTGACCGTGGTCCC
M-JH2	TGAGGAGACTGTGAGAGTGGTGCC
M-JH3	TGCAGAGACAGTGACCAGAGTCCC
M-JH4	TGAGGAGACGGTGACTGAGGTTCC

VL forward primer without restriction sites

M-VK1	GACATTGTGATGACACAGTCTCC
M-VK2	GATGTTGTGATGACCCAAACTCC
M-VK3	GATATCCAGATGACACAGACTCC
M-VK4	CAAATTGTTCTCACCCAGTCTCC
M-VL1	CAGGCTGTTGTGACTCAGGAATC

VL backward primers without restriction sites

M-JK1	TTTGATTTCCAGCTTGGTGCCTCC
M-JK2	TTTTATTTCCAGCTTGGTCCCCCC
M-JK3	TTTCAGCTCCAGCTTGGTCCCAGC
M-JL1	ACCTAGGACAGTGACCTTGGTTCC

Fig 9

VH forward primers with restriction sites

MVH1 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGCAGCTTCAGGAGTCAGG
MVH2 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCCAGGTGCAGCTGAAGGAGTCAGG
MVH3 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTCCAGCTGCAACAGTCTGG
MVH4 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTTCAGCTGCAGCAGTCTGG
MVH5 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCCAGGTCCAACTGCAGCAGCCTGG
MVH6 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCCAGGTTCAGCTGCAGCAGTCTGG
MVH7 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGAAGCTGGTGGAGTCTGG
MVH8 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGAAGCTGGTGGAATCTGG
MVH9 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTTCAGCTTCAGCAGTCTGG

VH backward primers with restriction sites

MJH1 Sall	GAGTCATTCTCGTGTCGACACGGTGACCGTGGTCCC
MJH2 Sall	GAGTCATTCTCGTGTCGACACTGTGAGAGTGGTGCC
MJH3 Sall	GAGTCATTCTCGTGTCGACACAGTGACCAGAGTCCC
MJH4 Sall	GAGTCATTCTCGTGTCGACACGGTGACTGAGGTTCC

VL forward primers with restriction sites

MVK1 ApaL1	TGAGCACACAGTGCACTCGACATTGTGATGACACAGTCTCC
MVK2 ApaL1	TGAGCACACAGTGCACTCGATGTTGTGATGACCCAAACTCC
MVK3 ApaL1	TGAGCACACAGTGCACTCGATATCCAGATGACACAGACTCC
MVK4 ApaL1	TGAGCACACAGTGCACTCCAAATTGTTCTCACCCAGTCTCC
MVL1 ApaL1	TGAGCACACAGTGCACTCCAGGCTGTTGTGACTCAGGAATC

VL backward primers with restriction sites

M-JK1 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTGATTTCCAGCTTGGTGCCTCC
M-JK2 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTTATTTCCAGCTTGGTCCCCCC
M-JK3 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTCAGCTCCAGCTTGGTCCCAGC
M-JL1 Not1	GAGTCATTCTCGACTTGCGGCCGCACCTAGGACAGTGACCTTGGTTCC

Fig. 9

Polypeptide or NPB	SEQ ID No.	Protein Sequence	SEQ ID No. of the CDR 3 as underlined
scFv1	1	1 EVQLQQSGPE LVKPGALVKI SCKASGYTVT SYDINWVKQR PGQGLEWIGW 51 IYPGDGSTKY NEKFKGKATL TVDKSSTTVY MQLSSLTSEN SAVYFCARGG 101 KYFDYWGQGT TLTVSTGGGG SGGGGSGGGG SALDIVMTQS PKFMSTSVGD 151 RVSVTCKASQ NVATNVAWYQ QKPGQSPKPL TYSASFRSSG VPDRFTGSGS 201 GTDFTLTISN VQSEDLAEYF CQQYNSYPYT FGGGTKLEIK AAAGAPVPYP 251 DPLEPRGAAS AWSHPQFEK*	73
scFv2	2	1 EVQLLESGGG LVQPGGSLRL SCAASGFTFS SYAMSWVRQA PGKGLEWVSA 51 ISGSGGSTYY ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARDS 101 GLQQGPRRRG ARVNFSYYGL DVWGRGTTVT VSSGGGGSGG GGSGGGGSAQ 151 AVLTQPSSAS GTPGQRVTIS CSGSNSNIGR NYVFWYQQFP GTAPKLLIYR 201 NNQRPSGVPD RFSGSKSGTS ASLAISGLRS EDEADYYCAS WDDSLTWVFG 251 GGTKVTVLGA AAGAPVPYPD PLEPRGAASA WSHPQFEK*	74
scFv3	5	1 ASVKVSCKTS GYTFIAYYIH WVRQAPGQGL EWMGRINPNT GGINLAQKFQ 51 GRVTVTRDTS ISTAHMELSR LSSDDTAVYY CARERIVPAG LRNRGMVTAV 101 GMDVWGRGTL VTVSSGGGGS GGGGSGGGGS AQSVVTQPPS MSGTPGQRVT 151 ISCSGSRSNI GRNYVYWYQQ FPGTAPKLLI YRNNERPSGV PDRFSASKSG 201 TSASLAISGL RSEDEADYYC ATWDDSLSGT WVFGGGTKLT VLGAAA	75
scFv4	6	1 LLESGGGLVQ PGGSLRLSCA ASGFTESSYA MSWVRQAPGK GLEWVSAISG 51 SGGGTYYADS VKGRFTISRD NSKNTLYLQM NSLRAEDTAV YYCARGGGRY 101 DSSHGEFDSWG RGTMVTVSSG GGGSGGGGSG GGSALSYEL TQPPSVSVAP 151 GETATITCGG RSLGSKVVHW YQQKPGQAPT LVIYYDSVRP SGVPERFSAS 201 NSRLSATLTV SRVEAGDEAD YYCQVWDRSS DHYVFGTGTK LTVLGAAA	76
scFv5	7	1 QLLESGGGLV QPGGSLRLSC AASGFTFSSY AMSWVRQAPG KGLEWVSAIS 51 GSGGSTYYAD SVKGRFTISR DNSKNTLYLQ MNSLRAEDTA VYYCARDWRW 101 QQFGGWEDPW GRGTLVTVSS GGGGSGGGGS GGGGSALETT LTQSPATLSL 151 SPGETATLFC RASQSVRNNL VWYQQKLGQA PRLLIFGAST RASGIPDRFT 201 GSGSGTDFSL TITKLEPEDE AVYYCQRYGG EPITFGQGTR LEIKRAAA	77
scFv6	8	1 QLVQSGGGLV QPGGSLRLAC EASGFRFSSY GMSWVRQAPG KGLEWVSSMS 51 DSGANTYYAD SVKGRFTISR DNAKKMLYLQ MSSLRGEDTA VYYCATLFRG 101 SGYVRHWGRG TLVTVSSGGG GSGGGGSGGG GSAQAVLTQP SSASGTPGQR	78

Fig. 40

	pos	sequence	No.
	151	VIISCSGSSS NIASNYVYWY QQLPGTAPKL LISKNSRRPS GVPDRFSGSK	
	201	SGTSASLAIS ELRSEDEADY YCAAWDDRLS GPAFGGGTKL TVLGAAA	
scFv7 (9)	1	KKPGSSVKVS CKASGGTFSS YAISWVRQAP GQGLEWMGGI IPMSGTPNYA	79
	51	QKFQDRVTIT ADKSTSTAYM ELSSLRSEDT AVYYCARGGR YVDFGRGPSY	
	101	HYYYMDVWGR GTLVTVSSGG GSGGGGSGG GSAQSVLTQ PPSASGTPGQ	
	151	RVTISCSGAT SNIGRNYVYW YHQLPGTAPK LLIYRNDQRP SGVPDRFSGS	
	201	KSGTSASLAI SGLRSDDEAD YYCAAWDDNL SGLFFGGGTK LTVLGAAA	
scFv8 (10)	1	AQVQLQQWGP GLVKASEILS LNCTVSGSSL SSGGYYWSWI RQHPGKGLEW	80
	51	IGYIHYSGST YYNPSLKSRV TISVDTSKNQ ESLKLSSVTA ADTAVYYCAR	
	101	VPLREDGEFDV WGQGTLVTVS SGGGGSGGGG SGGGGSDIQM TQSPSTLSAS	
	151	IGDRVTITCR ASEGIYHWLA WYQQKPGKAP KLLIYKASSL ASGAPSRFSG	
	201	SGSGTDFTLT ISSLQPDDEA TYYCQQYSNY PLTFGGGTKL EIKRAAA	
scFv9 (11)	1	ELKKPGSSVK VSCKAPRGTF NSYALNWVRQ APGQGLEWMG GIIPIFGSAN	81
	51	YAPKEQGRVT ITADESTTTA YLELSSLRSE DTAVYYCARA LHLDYVWRTY	
	101	NYYEDNWGKG TMVTVSSGGG GSGGGGSGGG GSALSSELTQ DPAVSVALGQ	
	151	TVRITCQGDS LRSYYASWYQ QKPGQAPVLV IYGKNSRPSG IPDRFSGSDS	
	201	GNTASLTITG AQAEDEADYY CNSRDRSGNR VVEGGGTKLT VLGAAA	
scFv10 (12)	1	SLRLSCAASG FTESSYAMSW VRQAPGKGLE WVSAISGSGG STYYADSVKG	82
	51	RFTISRDNSK NTLYLQMNSL RAEDTAVYYC ARGVTYHYDH DRRGVTAQIY	
	101	NHGLDVWGRG TTVTVSSGGG GSGGGGSGGG GSAQAVLTQP SSASGTPGQR	
	151	VTISCSGSSS NIGKNYVYWY QQLPGTAPKL LIYRNNQRPS GVPDRFSGSK	
	201	SGTSASLAIS GLRSEDEADY YCAARDNGLS AYVIFGGGTK LTVLGAAA	
scFv11 (13)	1	VKKPGESLKI SCKGSGYSFP NYWIAWVRQM PGKGLEWMGI IYPGDSDTIY	83
	51	SPSFRGQVTI SADKSISTAY LQWSSLKASD TAMYYCARQG CSGGKCYEKM	
	101	YASDIWGRGT LVTVSSGGGG SGGGGSGGGG SALSYELTQP PSASGTPGQR	
	151	VTISCSGSTS NIGRNSVFWH QQLPGTAPKV LISSDNQRPS GVSDRFSGSD	
	201	SGTSASLVIS RLRFEDEGDY YCAAWDDSLS AYVFGSGTKL TVLGAAA	
scFv12 (14)	1	AEVKKPGSSV RVSCKASGDT FSYNAINWVR QAPGQGLEWM GGIIPMFGTA	84
	51	KQAQKFQGRV TETADESTST AYMELTRLRS EDTAMYYCAR RGSYSNYERG	
	101	YYYHMDVWGQ GTLVTVSSGG GSGGGGSGG GSGSALPVLTQ PPSASGAPGQ	
	151	RITISCSGST ENIGRNYVDW YKQLPGTAPK LFIYKNDQRP SGVPDRFSGS	
	201	KSGTSASLVV SGLRSEDEAD YYCLTWDDSL SGPVFGGGTK LTVLGAAA	
scFv13 (15)	1	LQESGPGLVK PSGTLSLTCA VSGGSINNNN WWSWVRQPPG KGLEWIGEIY	85
	51	QSGSTNYNPS LKSRVTISVD KSNNQFSLKM SSVTAADTAV YYCARLNWNH	
	101	GPYYGMDVWG RGTLVTVSSG GGGSGGGGGS GGGSAQSVLT QPPSASGTPG	
	151	QRVTISCSGS SSNIGSNFVY WYQQLPGTAP KLLIYRNNQR PSGVPDRFSA	
	201	SKSGTSASLA ISGLRSEDEA DYYCAAWDDR RVVFGGGTKL TVLGAAA	

Fig. 10

		Sequence	
scFv14	1 51 101 151 201	VQLQESGPGL VKPSETLSLT CTVSGGPVAS SSYYWGFIRQ PPGKGLEWIG SIYDGGYTYY SPSLKSRATI SEDTSKNQVS LNLTSVTAAD TAVYYCAKDP GSLSAFWGQG TLVTVSSGGG GSGGGGSGGG GSALDIQLTQ SPSSLSASVG DRVTITCRTS QRISSYLNWY QQKPGKAPKL LIYAASSLQS GVPSRFSGSG SGTDFTLTIS SLQPEDFATY YCQQSYSTPI TFGQGTRLEI KRAAA	86
scFv15	1 51 101 151 201	LLESGGGLVQ PGGSLRLSCA ASGFTFSSYA MSWVRQAPGK GLEWVSAISG SGGSTYYADS VKGRFTISRD NSKNTLYLQM NSLRAEDTAV YYCARDWRWQ QFGGWEDPWG RGTLVTVSSG GGGSGGGGSG GGGSALDVVM TQSPATLSVS PGERVTLSCR ASQSVGSKLA WYQQKPGQAP RLLIFGTSTR ASGIPARESG SGSGTEFTLT ISSLQSEDEA VYYCQQYNNW PPYTFGQGTK VEIKRAAA	87
scFv16	1 51 101 151 201	AEVKKPGDSV KVSCKASGYR FETYGFSWVR QAPGQGLEWM GWINTYNGKT NYAQKFQGRV TMTTDTSTST AYMELRSLRS DDTAVYFCSR AEDDSRGYWN HYFSDYWGRG TTVTVSSGGG GSGGGGSGGG GSAQSVLTQP PSASGTPGQR VTISCSGSSS NIGSNYVYWY QQLPGTAPKL LIHKNNRRPS GVPDRFSGSK SGTSASLAIS GLRSEDEADY HCAAWDDSLS AVFGGGTKV TVLGAAA	88
scFv17	1 51 101 151 201	LESGGGLVQP GGSLRLSCAA SGFTFSSYAM SWVRQAPGKE LEWVSAISGS GGSTYYADSV KGRFTISRDN SKNTLYLQMN SLRAEDTAVY YCARDWRWQQ FGGWFDPWGR GTMVTVSSGG GGSGGGGSGG GGSALETTLT QSPGTLSLSP GDRATLSCRA SHSVSHNHLA WYQQNPGQAP RLLIFGASSR AAGIPDRFSG SGSGTDFTLT ISRLEPEDFA SYYCQQYGSP RRTFGQGTKV EIKRAAA	89
scFv18	1 51 101 151 201	KKPGSSVRVS CKAPGGTFGN SAISWVRQTP GQGLEWMGGI IPMFTTANYA QKFQGRVTIT ADKSTTAHM ELSSLRSEDT AVYYCARGGL GRFFDGPSHE SYYMEVWGKG TLVTVSSGGG GSGGGGSGGG GSAQSVLTQP PAASGTPGQR VTISCSGSNS NIGRNYVYWY QQLPGAAPKL LIYRNNQRPS GVPDRFSGSK SGPSASLAIS GLRSEDEADY YCAAWDDSLS GPAFGGGTKL TVLGAAA	90
scFv19	1 51 101 151 201	VQLVQSGAEV KKPGSSVKVS CKASGGTFSS DAISWVRQAP GQGLEWMGRI IPLINIPNYA QKFQGRVTIT ADKSTTTAYM ELTSLRFEDA AVYYCARVNN WNAEFDQWGRG TLVTVSSGGG GSGGGGSGGG GSALSSELTQ DPAVSVALGQ TVRITCQGDT LTSYYAAWYQ QKPGQAPLLV FYGKDKRPSG IPERFSGSSS GNIASLTITG AQAEDEGDFY CSSRDSSGYR EVFGAGTKLT VLGAAA	91
scFv20	1 51 101 151 201	KKPGSSVKVS CKASGGTFTS YAISWVRQAP GQGLEWMGGF IPVFGTANYA QKLQGRVTIT ADDSMTTVYM ELSSLTSEDT GVYYCARDLM RLARRDEYYY YYMDVWGQGT MVTVSSGGGG SGGGGSGGGG SAQSVLTQPP AASGTYGQKI TISCSGSSSN IGVNYVYWYR QFPGAAPHVV IYNNDQRPSG VPDRFSGSKS GTSASLAISG LRSEDEADYY CSTWDDTLSG YIFGVGTKVT VLGAAA	92
scFv21	1 51	QPGGSLRLSC AASGFTFSSY AMSWVRQAPG KGLEWVSAIS GSGGSTYYAD SVKGRFTISR DNSKNTLYLQ MNSLRAEDTA VYYCARDWRW QQFGGWFDPW	

Fig. 10

		Seq
	101 GQGTLVTVSS GGGGSGGGGS GGGGSALSSE LTQDPAVSVA LGQTVRITCQ 151 GDNLRSFSAS WYQLKPGQAP VLVIYGKNNR PSGIPDRFSA SSSGNTASLA 201 ITGALAEDEA DYYCNSRDSS GNPYVFGTGT KVTVLGAAA	93
scFv22 24	1 SSVKVSCKIS GGNLRLTVT WVRQAPGQGL EWVGRILPDS VNQVVKFQRR 51 LKLTSDTTSTR TAYLELRSLK SEDTAVYYCA ASSKIGFQVG ELDYWGRGTL 101 VTVSSGGGGS GGGGSGGGGS AQSVVTQPPS ASATPGQRVT ISCSGSSSNI 151 GRNYVYWYQQ VPGTAPQLIV YNNNQRPSGV PDRFSGSKSG TSASLGISGL 201 RSEDEADYYC STWDDSLSSP VEGGGTKLTV LGAAA	94
scFv23 25	1 TFSSYAMSWV RQAPGKGLEW VSAISGSGGS TYYADSVKGR FTISRDNSKN 51 TLYLQMNSLR AEDTAVYYCA RGRRRERSIN MIRGVRPQYD DSGMDVWGRG 101 TLVTVSSGGG GSGGGGSGGG GSALSYVLTQ PPSASGTPGH RVTISCSGSS 151 SNIGSNYVYW YQQLPGTAPK LLIYRNNQRP SGVPDRFSGS KSGTSASLAI 201 SGLRSEDEAD YYCAAWDDTL SGVLFGGGTK LTVLGAAA	95
scFv24 26	1 YAMSWVRQAP GKGLEWVSAI SGSGGSTYYA DSVKGRFTIS RDNSKNTLYL 51 QMNSLRAEDT AVYYCARNTG KGITLVRGVY CQDCDRSSTS RMDVWGQGTL 101 VTVSSGGGGS GGGGSGGGGS AQAVLTQPSS ASGTPGQRVT ISCSGSTSNI 151 GRNYVDWYQQ LPGTAPKLLI YRNNKRPSGV PDRFSGSKSG TSASLAISGL 201 RSEDEADYYC AAWDDSLSGW VFGGGTKLTV LGAAA	96
scFv25 27	1 GLVQPGGSPR LSCAASGFTF SSYAMSWVRQ APGKGLEWVS AISGSGGSTY 51 YADSVKGRFT ISRDNSKNTL YLQMNSLRAE DTAVYYCAKD MGYSYGYGTR 101 GLFDYWGRGT MVTVSSGGGG SGGGGSGGGG SAQSVVTQPP SASGAPGQRI 151 TISCSGSTFN IGRNYVDWYK QLPGTAPKLF IYKNDQRPSG VPDRFSGSKS 201 GTSASLVVSG LRSEDEADYY CLTWDDSLSG PVFGGGTKVT VLGAAA	97
scFv26 28	1 ESGGGLVQPG GSLRLSCAAS GFTFSSYAMS WVRQAPRKGL EWVSAISGSG 51 GSTYYADSVK GRFTISRDNS KNTLYLQMNS LRAEDTAVYY CARDWRWQQE 101 GGWFDPWGRG TTVTVSSGGG GSGGGGGGSA GSALETTLIQ SPATLSVSPG 151 DRATLSCRAS QSIGGNLAWY QQKPGQPPRL LIFGASTRAS GTPARFSGSG 201 SGTEFTLTIS SLQSEDFAVY YCQQYNNWPP WTFGQGTRLE IKRAAA	98
scFv27 29	1 QPGGSLRLSC AASGFTFSSY AMSWVRQAPG KGLEWVSAIS GSGGSTYYAD 51 SVKGRFTISR DNSKNTLYLQ MNSLRAEDTA VYYCAKGDGV VAGTTYYYG 101 MDVWGRGTTV TVSSGGGGSG GGGGGGSA QSVLTQPPSA SGAPGQRITI 151 SCSGSTFNIG RNYVDWYKQL PGTAPKLFIY KNDQRPSGVP DRFSGSKSGT 201 SASLVVSGLR SEDEADYYCL TWDDSLSGPV FGGGTKLTVL GAAA	99
scFv28 30	1 ASGFGLNGYE MHWVRQAPGQ RLEWLGRINA AIGDTRYSRE FQDRVSITRD 51 MSANTVYMEM SRLRFEDTAV YYCVRFHDWR HCNSATCQPP EDHWGKGTLV 101 TVSSGGGGSG GGGSGGGGSA LSSELTQDPA VSVALGQTVR ITCQGDSLRY 151 YSASWYRQKP GQAPVIVMYG NTRRPSGIPD RISGSSSGNT ASLTISGAQA	100

Fig. 10

	201	EDEADYYCNS RDSSGNHLVF GGGTKLTVLG AAA	
scFv29	1	VQPGGSLRLS CAASGFTFSS YAMSWVRQAP GKGLEWVSAI SGSGGSTYYA	101
	51	DSVKGRFTIS RDNSKNTLYL QMNSLRAEDT AVYYCARDHR SGRGGGSYLL	
	101	RPLDYWGQGT MVTVSSGGGG SGGGGSGGGG SALPVLTQPP SASGTPGQRV	
	151	TISCSGSSSN IGRNYVYWYQ QLPGTAPKLL IYRNNLRPSG VPDRFSGSKS	
	201	GTSASLAISG LRSEDEADYY CAAWDDTLSG VVFGGGTKLT VLGAAA	
scFv30	1	EVRKPGASVK ISCKASGFTF TSYLFHWVRQ APGQRLEWMG WINAGNGNTK	102
	51	YSPKFQGRVT LTGDTSTSTT YMELSSLTSE DTAVYYCARD QVFYESGSYY	
	101	IRPSEDFWGR GTLVTVSSGG GSGGGGSGGG GGSDIQMTQS PSTLSASIGD	
	151	RVTITCRASE GIYHWLAWYQ QKPGKAPKLL IYKASSLASG APSRFSGSGS	
	201	GTDFTLTISS LQPDDEATYY CQQYSNYPLT EGGGTKLEIK RAAA	
scFv31	1	VRPGGSLRLS CAASGFTEDD YGMSWVRQAP GKGLEWVSGI NWNGGSTGYA	103
	51	DSVKGRFTIS RDNAKNSLYL QINSLRAEDT AVYYCARRRY ALDYWGRGTM	
	101	VTVSSGGGGS GGGGSGGGGS ALSSELTQDP ATVSVALGQT VRITCQGDSL	
	151	DKYYATWYQQ KPGQAPLLVF FSENRRPSGI PDRFSGSNSG NTASLTITGA	
	201	QAEDEADYYC NSREIGTNRI LEGGGTKLTV LGAAA	
scFv32	1	LVQPGGSLRL SCAAAGFTFS TFEMNWVRQA PGKGLEWVSY ISGSGHAIYY	104
	51	ADSVKGRFTI SRDNANNSLY LQMNSLTAED TAVYYCAREK YQLLLGKYDY	
	101	GMDVWGRGTT VTVSSGGGGS GGGGSGGGGS ALPVLTQPPS ASGTPGQRVT	
	151	ISCSGSSSNI GSNTLNWYQQ LPGTAPKLLI YSNDQRPSGV PDRFSGSKSG	
	201	TSASLAISGL QSEDEADYYC AAWDDSLNGW VFGGGTKVTV LGAAA	
scFv33	1	RASGGTSSSS AFSWVRQAPG QGLQWMGGII PLFGAANYAQ KVRAGLTITA	105
	51	DESTGTSYMK LENLQSDDTA VYFCATNGQT RSPPGYYYGM DVWGRGTLVT	
	101	VSSGGGGSGG GGSGGGGSAQ SVLTQLPSAS GAPGQRITIS CSGSTENIGR	
	151	NYVDWYKQLP GTAPKLEIYK NDQRPSGVPG RESGSKSGTS ASLVVSGLRS	
	201	EDEADYYCLT WDDSLSGPVF GGGTKLTVLG AAA	
scFv34	1	ACKGFGYTEV DHGISWVRQA PGQGLEWMGW INTHDGHTNY AQKTQARLTM	106
	51	TTDASINTSY MELRSLTSDD TAVYYCARGG ETRTAHRSRR ATNDNGYPYY	
	101	SSGLDVWGQG TLVTVSSGGG GSGGGGSGGG GSAQAVLTQP SSASGTPGQR	
	151	VTISCSGSSS NIGSNYVYWY QQLPGTAPKL LIYRNNQRPS GVPDRFSGSK	
	201	SGTSASLAIS GLRSEDEADY YCAAWDDSLS GWVFGGGTKL TVLGAAA	
scFv35	1	PGASVKVSCK ASGYTFTSYY MHWVRQAPGQ GLEWMGIINP SGGSTSYAQK	107
	51	FQGRVTMTRD TSTSTVYMEL SSLRSEDTAV YYCARGSGAR MVRGVIIDPY	
	101	GMDVWGRGTL VTVSSGGGGS GGGGSGGGGS AQSVLTQPPS ASGTPGQRVT	
	151	ISCSGSSSNV GSNYVSWYQQ FPGTAPKLLI YRNNQRPSGV PDRFSGSKSG	
	201	ISASLAISGL RSEDEADFYC VAWDDSLREY VFGTGTKVTV LGAAA	

Fig. 10

scFv36	38		108
		1 ESGGGLVQPG GSLRLSCAAS GFTFSSYAMS WVRQAPGKGL EWVSAISGSG 51 GSTYYADSVK GRFTISRDNS KNTLYLQMNS LRAEDTAVYY CAKGGTRVTH 101 RGGFDIWGRG TMVTVSSGGG GSGGGGSGGG GSALPVLTQP PSASGAPGQR 151 ITISCSGSTF NIGRNYVDWY KQLPGTAPKL FIYKNDQRPS GVPDRFSGSK 201 SGTSASLVVS GLRSEDEADY YCLTWDDSLS GPVEGGGTKL TVLGAAA	

Fig. 10

Ref.:	SEQ ID No.	Nucleotide Sequence	
scFv1	3	1	GAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTTAGTGAAGATATCCTGCAAGG
		71	CCTCGGGATACACCGTCACAAGCTACGATATAAACTGGGTGAAGCAGAGGCCTGGACAGGGACTTGAGTG
		141	GATTGGATGGATTATCCTGGAGATGGTAGTACTAAGTACAATGAGAAGTTCAAGGGCAAGGCCACACTG
		211	ACTGTAGACAAATCCTCCACCACAGTCTACATGCAGCTCAGCAGCCTGACTTCTGAGAACTCTGCAGTCT
		281	ATTTCTGCAAGAGGTGGTAAATACTTTGACTACTGGGGCCAAGGCACCACTCTCACAGTGTCGACAGG
		351	TGGAGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACTGCACCTGCAAGGCCAGTCAGAAGTGGCCTA
		421	CCAAAATTCATGTCCAATCAGTAGGAGACAGGGTCAGCGTCACCTGCAAGGCCAGTCAGAATGGCCTA
		491	CTAATGTAGCCTGGTATCAACAGAAACCAGGGCAATCTCCTAAACCACTGACTTACTCGGCATCCTTCCG
		561	GTCCAGTGGAGTCCCTGATCGCTTCACAGGCAGTGGATCTGGACAATATAACAGCTATCCGTACACGTTCGGAGGGG
		631	GTGCAGTCTGAAGACTTGGCAGAGTATTTCTGTCAGCAATATAACAGCTATCCGTACACGTTCGGAGGGG
		701	GGACCAAGCTGGAAATAAAAGCGGGCCCAGTGCGCCGAGTCCGTATCCAGATCCGCTGGAACCGCGTGG
		771	GCCCGCAAGCGCTTGGAGCCACCCGCAGTTCGAAAAATAA
scFv2	4	1	GAGGTGCAGCTGTTGGAGTCTGGGGGAGGCTTGGTGCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAG
		71	CCTCTGGATTCACCTTTAGTCAGCTATGCCATGGGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTG
		141	GGTCTCAGCTATTAGTGGTAGTGGTGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATC
		211	TCCAGAGACAATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGT
		281	ATTACTGTGCGCGAGACTCGGGGTGGGGGGGACCACGGGTCACCGTCTCCTCAGGTGGAGGCGGTTCAGGCGGA
		351	CTACGGTCTGGACGTCTGGGGGTGGCGGAAGTGCACTGCACCTGCACCTGTCGCTGTGCTGGTTGGGGACCCCCG
		421	GGTGGCTCTGGCGGTGGCGGAAGTGCACTGCACCTGCAGCCGTCTCCAGCCTCTGGGGACCCCCG
		491	GGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAACTCCATCTACAGGAACAATCAGCGGCCCTCAGGGGTACCA
		561	GCAGTTCCCAGGAACGGCCCCCCAAAATCTCGCACATCAGCCTCCCGGTCCGAGGGGTCCCTGAC
		631	CGATTCTCTGGCTCCAAGTCTGGCACATGGGATGACAGCGCCTGGTGTTCGGCGGAGGACCAAGGTCACCGT
		701	CCTAGGTGCGGCGCCGCAGGTGCGCGGTGTGTATCCAGATCCGCTGGAACCGCGTGGGGGCGCCAAGCGCT
		771	TGGAGCCACCCGCAGTTCGAAAAATAA
scFv3	39	1	GGGCCCTCAGTGAAGGTCTCCTGCAAGACCTTCATCGCCTATTATATTCATTGGGTGCG
		71	ACAGGCCCCTGACAAGGGGCTTGAGTGGATGGGACGGATCAACCCTAACACTGGTGGCATAAACCTTGCA
		141	CAGAAGTTTCAGGGCAGGGTCACCGTCACCGTGACCAGGGACGACGTCCATCAGCACACAGCCCACATGAGCTGAGTA
		211	GGCTGAGCTCTGACGACACGGCCGTGTTGGGCCTATACTACTGTGCGAGAGAGATCGTACCAGCTGGTCTGAGGAA
		281	CCGTGGGATGGTGACTGCGGTTGAATGGACGTCTGGGGCCGGGAACCCTGGTCACCGTCTCGAGTGGA
		351	GGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACAGTCTGTCTGACGCAGCCGCCCT
		421	CAATGTCTGGGACCCCCGGGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAGTTCCAACATTGGAAGGAA
		491	TTATGTATACTGGTACCAGCAGTTCCCAGGAACGGCGCCCCAAAGTCTGCACCTCAGCCTCCTCAGCCTCCTCAGCCTCAGCCTCAGTAATGAACGG
		561	CCCTCAGGGGTCCCTGACCGATTCTCTGCCTCCAGTGTCTGGCACCTCAGCCTCCCCATCAGTGGAC

Fig. 11

	Pos	Sequence
	631	TCCGGTCCGAGGATGAGGCTGATTATTACTGCGCAACGTGGGATGCAGTCTGAGTGGGACTTGGGTGTT
	701	CGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv4 40	1	CTGTTGGAGTCTGGGGGAGGCTTGGTGCTACAGCTGGGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTG
	71	TCACCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGC
	141	TATTAGTGGTAGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGAC
	211	AATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTG
	281	CGAGAGGTGGGGGCGGCGGTCGGATATGATAGTAGTAGTACTGGTTGGGGGCGGGGGACAATGGTCACCGT
	351	CTCGAGTGAGGGCCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACTTCCTATGAGCTG
	421	ACTCAGCCACCCTCAGTGTCTGGTGCCCCAGGAGAGACGGCCACCAATTACCTGTGGGGGACGCAGCCTTG
	491	GCTCCAAAGTTGTCATTGGTATCAGCAGGACGGCTCAGCCCCTACATTCTCGGTTCATTTATTATGATTC
	561	CGTCCGGCCCCTCGGGGGTCCCTGAGCCGATTCTCTGCCTCCAACTCTCGGTTGTCGGGGCCCACCCTGACCCGTC
	631	AGCAGGGTCGAAGCCCGGGGATGAGGCCGGGATGAGGCGGATAGAAGTAGTGACCATTATG
	701	TCTTCGGAACTGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv5 41	1	CAGCTGTTGGAGTCTCGGGGGAGGCTTGGTACAGCCTGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTG
	71	GATTCACCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTC
	141	AGCTATTAGTGGTAGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGA
	211	GACAATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACT
	281	GTGCCGAGAGATTGGAGATGGCCAACAATTGGGGGCGTGGTCTCAGGGGAGGTGGCTCTGGCGGTGGCGGAAGTCCGTGGTCAC
	351	CGTCTCGAGTGCAGTCTCCTGCCACCCTGTCTCTGTCTCCAGGGGAAAACAGCCACCCTCTTCTGCAGGGCCAGTC
	421	CTCACGCCAGTCTCCTGCCACCCTGTCTGTCTGGTACCAGCAGAAGCCAGGCCAGGTCCCTCCATCTTTGG
	491	AGAGTGTTAGGGAACAACTTAGTCTGGTACCAGACAGGTTCACTGGCAGTGGGTCTGGGACACTCCTCAGTCTC
	561	TGCATCCACCAGGGCCAGTGGCATCCCAGGATTTGCAGTGTATTACTGGCAGTGTATTACTGTCAGCGGTATGGTGGTTTCCCGATCA
	631	ACCATCACCAAAACTGGAGCCTGAGGCTGAGGACTGGAGATTAAAACGTGCGGGCGC
	701	CCTTCGGCCGCCAAGGAGGACACGACGCTTGGTGCTGGAGCTTGGTGTGCAGC
scFv6 42	1	CAGCTGGTGCAGTCTGGGGGAGGCTTGGTGCAGCCTGGGGGGTCCCTGAGACTCGCCTGTGAAGCCTCTG
	71	GATTCAGGTTTAGCAGCTATGGCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTTGAGTGGGTCTC
	141	TTCCATGAGCGATAGTGGTGCCAATACATACTACGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGA
	211	GACAATGCCAAGAACAGATGTTGTATCTACAAATGAGCAGCCTGAGAGCCGAGGACACGGCCGTGTATTACT
	281	GTGCGACTCTCTTTAGGGGAGTGGCTACGTTCGTCTACTGCGGTCTGGGCTGGGGGCCGAGGGAACCCTGGTCGTCTCGAG
	351	TGGAGGCCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACAGGGTCGTGCTGACTCAGCCG
	421	TCCTCAGCCGTCTGGGACCCCGGGACAGTCATCATCTCTTGTTCTGGAAGCAGCTCCAACATCGCAA
	491	GTAATTATGTATACTGGTACCAGCAACTCCCAGAAGCGCCCCAAACTCCTCATCTCTAAGAATAGTCG
	561	GCGCCCTCCGGTCCAGGGTTCCTGAGGATGAGGCTGATGTGCAGCATGGGATGACAGCCTGAGTGGTCCGGCAT
	631	GAACTCCGGTCCGAGGATGAGGCTGATTATTACTGCAGGCTGACCGTCCTAGGTGCGGGCGC
	701	TCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv7 43	1	AAGAAGCCTGGGTCTCCGGTGAAGTCTTCCTGCAAGGCTTCTGGAGGCACCTTCAGCAGTTATGCTATTA
	71	GTTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGGATGGGAGGGATCATTCCTATGTGCTGGTACACC

Fig. 11

	141	AAACTACGCACAGAGAGTTCCAGGACAGCAGAGTCACGATTACCGCGGACAAATCCACGAGCACAGCCTACATG
	211	GAGCTGAGCAGCCTGAGATCTGAGGACACGGCCGTGTATTACTGTGCCGAGGGGGGGCCTACGTTGACT
	281	TCGGTCGTGGCCCTTCGTACCACTACTACATGGACGTCTGGGCGGTGGCCAGGGGAACCCTGGTCACCGTCTC
	351	GAGTGGAGGCCGGCGGTTCAGGCGGACCCCCGGACAGAGGGTGGCTCACCATCTCTGTGGGCGACCATCTCTTGTTCTGGGGCCACCTCCAACATCG
	421	CCGCCCTCAGCGTCTGGGACACAGAGGGTCACCAACTCCCAGGACGCGCCCCAAGTCTCCTCATCTATAGGAATGA
	491	GAAGGAATTATGTTTACTGGTACCACCAACTCCCAGGACGCGCCCCAAGTCTCCTCATCTATAGGAATGA
	561	TCAGCGTCCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCCCTGGCCATC
	631	AGTGGCCTCCGGTCCGACGATGAGGCTGATTATTACTGTGCCGGTGGGACGACAACCTGAGTGGTCTAT
	701	TTTTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv8	44	1	GCCCAGGTGCAGCTACAGCCAGTGGGGCCCCAGACTGGTGAAGGCTTCGGAGATCCTGTCCCTCAACTGCA
	71	CTGTCTCTGGTGGTGGCTCCCTCAGCAGTGGTTACTACTGGAGCTGGATCCGCCAGCACCCAGGGAAGGG
	141	CCTGGAGTGGATTGGGTACATCCATTACAGTGGGAGCACGTACTACAACCCGTCCCTCAAGAGTCGAGTT
	211	ACCATATCAGTAGACACGTCTAAGAACCAGTTCTCCCTGAAGCTGAGCTCTGTGACTGCCGCGGACACGG
	281	CTGTGTATTATTGTGCGAGAGTTCCGTTGAGATTTGATGGTTTGATGTCTCGGGGCCAAGGCACCCTGGT
	351	CACCGTCTCGAGTGGTGGAGGCGGTTCAGGCGGAGGTGGCAGACAGAGTCACCATCACCTGCGGGCCAGTGAGG
	421	ACCAGTCTCCTTCCACCCTGTCTGTGGTATCTATTGGAGACAGAGTCACCATCACCTGCGGGCCAGTGAGG
	491	GTATTATCACTGGTTGGCCTGGTATCAGCAGAAGCCAGGGAAAGCCCCTAAACTCCTGATCTATAAGGAC
	561	CTCTAGTTTAGCCAGTGGGCCCCATCAAGGTTCAGCCGGCAGTGGATCTGGGACAGATTTCACTCTCACC
	631	ATCAGCAGCCTGCAGCCTGATGATTTTGCAACTTATTACTGCCAACAATATTAATTATCCGCTCACTT
	701	TCGGCGGAGGGACCAAGCTGGAGATCAAACGTGCGGCCGC
scFv9	45	1	CTGAGCTGAAGAAGCCTGGGTCCTCGGTAAAGGTCTCCTGCAAGGCTCCTGCAAGGCTCCTCCTGCAAGGCTCACCTTCAACAGTTA
	71	TGCTCTCAACTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGAGGGATCATCCCTATTTT
	141	GGTAGTGCAAATTACGCACAGAAGTTCCAGGGCAGAGTCACCATTACCGCGGACGAATCCACGACCACAG
	211	CCTACTTGGAGCTGAGCAGCCTGAGATCTGAGGACACGGCCGTGTATTACTGTGCGGAGCTCTCCATTT
	281	GGATTACGTTTGGAGGACTTATAATTACTTTGACAACTGGGCAAAGGACAATGGTCACCGTCTCG
	351	AGTGGAGGCGGCGGTTCAGGCGGAGGTGGCCTTGGGACAGCAGTCAGATCACATGCCAGGAGACAGCCTCAGAAG
	421	AGGACCCCTGTGTCTGTGACCCAGGAAGCCAGGACAGCCCCTGTCCTTGTCATCTATGGTAAAAATAGT
	491	TTATTATGCAAGCTGGTACCAGCAGAAGCCAGGAAACACAGCTTCCTTGACCATCACTG
	561	CGGCCCTCAGGGATCCCAGACCGATTCCTGGCTCCGACTACTGTAACTCCCGGGACAGAGAAGTGGTAACCGCGTGGTCTT
	631	GGGCTCAGGCGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
	701	CGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv10	46	1	TCCCTGAGACTCTCCTGTGCGCCCTCTGGATTCACCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGG
	71	CTCCAGGGAAGGGCCTGGAGTGGGTCTCAGCTATTAGTGGTAGTGGTAGCACATACTACGCAGACTC
	141	CGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTG
	211	AGAGCCGAGGACACGGCCGTGTATTACTGTGCGAGAGGGGTTACGTTACTACTATGACCATGACCATGACCAGGCGGTG
	281	GTGTGACCGCGCAAATATATAAACCACGGTTTGGAGCGTCTGGGGAGGGGACCACGGTCACCGTCTCGAG

Fig. 11

	351	TGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACAGGCTGTGCTGACTCAGCCG
	421	TCCTCAGCGTCTGGGACCCCCGGGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAGCAGCATCGGAA
	491	AGAATTATGTATACTGGTATCAGCAGCTCCCAGGAACGGCCCCCAAACTCCTCAGCCTCCCATAGGAATAATCA
	561	GCGGCCCTCAGGAGTCCCTGACCGATTCTCTGGCTCCGCAAGTCTGGCACTCCCAGTCCCATCAGT
	631	GGGCTCCGGTCCGAGGATGAGGCTGATTATTATTGTGCGGCACGGGATAACGGCCTGAGTGCTTATGTGA
	701	TATTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv11 47	1	AGGTGAAAAAGCCCGGGAGTCTCTGAAGATCTCCTGCAAGGGTTCTGGATACAGCTTTCCAACTACTG
	71	GATCGCCTGGGTGCGCCAGATGCCCGGGAAAAGCCTGGGAGTGGGGATCATCTATCCTGGTGACTCT
	141	GATACTATATACAGCCCGTCCTTCCGAGGCCAGGTCACCATCTCAGCCGACAAGTCCATCAGCACCGCCT
	211	ACCTGCAGTGGAGCAGCCTGAAGGCCTCGAGTGATTACTGTGCGAGAGACAGGGTTGTAGTGG
	281	TGGTAAATGCTACGAGAAAATGTATGCTTCTGATATCTGGGGCCAGGGAACCCTGGTCACCGTCTCGAGT
	351	GGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGCAGTGCACTTTCCTATGAGCTGACTCAGC
	421	CACCCTCAGCGTCTGGGACCCCCGGGCAGAGGGTCACCAGTTACCAGAACGGCCCCATCTCATCTTCTGATAAT
	491	AAGGAATTCTGTATTTGGCACCAGCAGTTACCAGGAACGGCCCCCAAAGTCCTCATCTCTTCTGATAAT
	561	CAGCGACCCTCAGGGGTCTCTGAAGATGGGGTTCTGGCTCCGCAAGTCTCGGCACCTCAGCCTCGTCATCA
	631	GTCGCCTCCGGTCCGAGGATGAGGGTGATTACTACTGTGCAGCAGTGGGATGACAGTCTGAGTGCTTATGT
	701	CTTCGGAAGTGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv12 48	1	GGGCTGAGGTGAAGAAGCCTGGGTCCTCGTGAGGGTCTCCTGCAAGGCTTCTGGAGACACCTTCAGCTA
	71	CAATGCTATCAACTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGGGATCATCCCTATG
	141	TTTGGTACAGCAAAGCAGGCACAGAAGTTCCAGGGCAGAGTCACGTTTACCGCGGAATCCACGAGCA
	211	CAGCCTACATGGAGTTGACTAGGGTGAGATCCGAGGACACGGCCATGTATTACTGTGCGGACGGGGCTC
	281	GTACAGTAATTACGAGAGGGGTATTACTATCACATGGACGTCTGGGGCCAGGGAACCCTGGTCACCGTC
	351	TCGAGTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACTGCCTGTGCTGACTC
	421	AGCCACCCTCAGCGTCTGGGACCCCCGGGCAGAGGGATCACCATCTCTTGTTCCGGAAGCACCTTCAACAT
	491	CGGGAGAAATTATGTTGACTGGTATAAACAACTCCCGGAACGGCCCCTAAACTCTTCATCTATAAGAAT
	561	GATCAGCGACCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCGCAAGTCTGACTGGGATGAGCCTGGTCG
	631	TAAGTGGACTCCGCTCCGAGGATGAGGGTGATTATTACTGTGCGGGATGGGATGACAGCCTGAGTGGTCC
	701	GGTGTTCGGGGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv13 49	1	GCTCAGGAGAGTCCGCCCAGGACTGGTGAAGCCTTCGGGGAGACCCTGTCCCTCACCTGCGCTGTCTCTGGT
	71	GGCTCCATCAACAATAATAATTGGTGGAGTTGGGTCCGCCAGCCCCCAGGGAAGGGGCTGGAGTGGATTG
	141	GGGAAATCTATCAGAGTGGGAGCACCAACTACAACCCGTCCCTCAAGAGTCGAGTCACCATATCAGTAGA
	211	CAAGTCCAACAACCAGTTCTCCCTGAAGATGAGTTCTGTGACCGCCGCGACACGGCCGTGTATTACTGT
	281	GCGAGGCTTAACTGGAACCACGGGCCCTACTACGGTATGGACGTCTGGGGCCGTGGCACCCTGGTCACCG
	351	TCTCGAGTGGAGGCGGCGGTTCAGGCCGTATGGAGGTGCTCTGGCGGTGGCGGAAGTGCACAGTCTGTGCTGAC
	421	GCAGCCGCCCTCAGCGTCTGGGACCCCCGGACAGAGAGTCCCCAGGAACGGCCCCCAAAACTCCTCATCTATAGA
	491	ATCGGAAGTAATTTGTATACTGGTACCAGCAGCTCCCTGACCGATTCTCTGGCTCCGCCATTCTCAGCCTCCCTGGC
	561	ATAATCAGCGGCCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCGCCATTCTCAGCCTCCCTGGC

	631	CATCAGTGGGCTCCGGTCCGAGGATGAGGCTGATTATTACTGTGCGGCATGGGATGACAGGCGTGTGGTA
	701	TTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv14	50	
	1	GGTGCAGCTGCAGGAGTCGGGCCCAGGACTGGTGAAGCCTTCGGAGACCCTGTCCCTCACCTGCACTGTC
	71	TCTGGCGGCGCCCCTGCTCCGCCAGTAGTAGTTACTACTGGGGCTTCATCCGCCAGCCCCCAGGAAAGGGCTGG
	141	AGTGGATTGGGGAGTATTTATGATGGTGGCTACACCAGTCTCCCTGAACCTGACCTGGGCCGCCTTTACCAT
	211	ATCCTTCGACACGTCCAAGAACCAGGTCTCCCTGAACCTGAGCTCTGTTACCGCCGCGGACACGGCCGTC
	281	TATTACTGTGCGAAAGACCCGGGCAGTTTGAGCGCGTTGGCGGCGGAAGTGCACTTGACCGTCTCGA
	351	GTGGAGCGCCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGGACATCCAGTTGACCCA
	421	GTCTCCATCCTCCCTGTCTGCCTGTCTGTGTAGGAGACAGAGTCACCATCACTTGCCGCGACAAGTCAGCGCATT
	491	AGCAGCTATTTAAATTGGTATCAGCAGAAGAAGCCAGGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCA
	561	GTTTGCAAAGTGGGGTCCCATCAAGGTTCAGTGGCCAGTGGTCTGGGACAGATTTCACTCTCACCATCAG
	631	CAGTCTGCAACCTGAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGC
	701	CAAGGGACACGACTGGAGATTAAACGTGCGGCCGC
scFv15	51	
	1	CTGTTGGAGTCTGGGGGAGGCTTGGTACAGCCTGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGAT
	71	TCACCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGC
	141	TATTAGTGGTAGTGGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGAC
	211	AATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTG
	281	CGAGAGATTGGAGATGGCAACAATTGGGGGCTCTGGGGCCAGGGAACCCTGGTCACCGTCTCGGTCACCGT
	351	CTCGAGTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGGACATTGTGATG
	421	ACTCAGTCTCCAGCCACCCTGTCTGTCTGGTCCTACCAGGGGAAAGAGTCACCCTCTCCTGCAGGGCCAGTCAGA
	491	GTGTTGGCAGCAAGTTGGCCTGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTTTGGTAC
	561	ATCAGCCAGGCCAGTGGCCAGTCTGCAGTTTGCAGTTTATTACTGTCAGCAGTATAATAACTGGCCTCCACC
	631	ATCAGCAGCTGCAGTCTGAAGATTTTGCAGTTTATTACTGTCAGCAGTATAATAACTGGCCTCCGTACA
	701	CTTTTGGCCAGGGGACCAAGGTGGAAATCAAACGTGCGGCCGC
scFv16	52	
	1	GCTGAGGTGAAGAAGCCTGGGGACTCAGTGAAGGTCTCCTGCAAGGCCTCTGGTTACAGGTTTGAAACCT
	71	ATGGTTTCAGCTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGATCAACACTTACAA
	141	TGGTAAGACAAATTATGCACAGAAGTTCCAGGGCAGAGTCACCATGACCACAGACACGTCCACGAGCACA
	211	GCCTACATGGAGTTGAGGAGCCTGAGATCGGACGACACGGCCGTGTATTTTGTTCGAGAGCTGAGGATG
	281	ATAGCAGAGGTTATTGGAACCATTACTTCTCCGACTACTGGGGGCGGGGGAAGTGCACCGTCTGCTGCTCGAG
	351	TGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGCACAGTCGTGCTGACTCAGCCA
	421	CCCTCAGCGTCTGGGACCCCCGGGCAGAGGGTCACCATCTGTTCTGGAAGCAGCTCCAACATCGGAA
	491	GTAATTATGTATACTGGTACCAGCAGCTCCCAGGAACGGCCCCCAAACTCCTCATCCATAAGAATAATCG
	561	GGGCCCTCCGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGGCTCTGACTCCATCAGT
	631	GGGCTCCGGTCCGAGGATGAGGCTGATTATCACTGTGCAGCGTGTGGAGATGACAGCCTGAGTGCTGTGTT
	701	TCGGCGGAGGGACCAAGGTCACCGTCCTAGGTGCGGCCGC
scFv17	53	
	1	TTGGAGTCTGGGGGAGGCTTGGTACAGCCTGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGATTCA
	71	CCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGAGAGCTGGGTCTCAGCTAT

	141 TAGTGGTAGTGGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAAT
	211 TCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGCGA
	281 GAGATTGGAGATGGCAACAATTGGGGGCTGGTTCGACCCTGGGGCCGGGGAACCATGGTCACCGTCTC
	351 GAGTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGGAAGTGCACTTGAAACGACACTCACG
	421 CAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGTGATAGAGCCACCCTCTCCTGCAGGGCCAGTCACAGTG
	491 TTAGTCACAACCACTTAGCCTGGTACCAGCAGAAATCCTGGCCAGGCTCCCAGGCTCCTCATTTTGGTGC
	561 ATCCAGCAGGGCCGCTGGCATCCCTGACAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACC
	631 ATCAGCAGACTGGAGCCTGAAGATTTTGCATCATATTACTGTCAGCAGTATGGTAGCCCCGGGCGGGACGT
	701 TCGGCCAAGGGACCAAGGTGGAAATCAAACGTGCGGCCGC
scFv18 54	1 GAAGAAGCCTGGGTCCTCGGTGAAGGCTCCTGGAGGCACCTTCGGCAACTCTGCTATC
	71 AGTTGGGTGCGACAGCCCCTGGACAAGGGCTTGAGTGGATGGGAGAATCATTCCTATGTTTACTACAG
	141 CAAACTACGCCACAGAAGTTCCAGGGCAGAGTCACCATTACCGCGGACAAATCCACGAGCACAGCCCACAT
	211 GGAGCTGAGCAGCCTGAGATCTGAGGACACGGCCGTCTATTACTGTGCGAGAGGCGGACTGGGACGATTT
	281 TTTGACGGCCCCTCCCACTTCTCCTACTACATGGAAGTCTGGGCGAAAGGAACCCTGGTCACCGTCTCGA
	351 GTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGCACCATCTCTGTTCTCGGAAGCAACATCGGA
	421 GCCCGCAGCGTCTGGGACCCCCGGGCAGAGGGTCCCAGGAGCGCTCCCAGGAGCAACTCCATCTATAGGAATAATC
	491 AGAAAATTATGTCTACTGGTATCAGCAGCTCCCTGGCTCGATTCCTGATTATTACTGGTCACCGTCCCCATCAG
	561 AGCCGGCCCCTCAGGGTCCGGAGGATGAGGCTGAATTACTGTGCAGCATGGGATGACAGCAGCCTGAGTGGCCCCTGCA
	631 TGGGCTTCCGGTCCGGAGGACCAAGCTGACCGTCCCTAGGTGCGGCCGC
	701 TTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv19 55	1 AGGTCCAGCTGGTACAGTCTGGGGCTGAGGTGAAGAAGCCTGGGTCGGTGAAGGTCTCCTGCAAGGC
	71 TCTGGAGGCACCTTCAGCAGCGATGCTATCAGCTGGGTGCGACAGGCCCCTGGACAAGGGACTTGAGTGG
	141 ATGGGAAGGATCATCCCTCTAATTAATATACCAAAACTACGCACAAACTACGAGCTGACAGAGTTCACGATTA
	211 CCGCCGGACAAATCCACGACCACACTACGCCTAAGATTGAGGACCCTGGTCCCGGGAACCCTGGTCACCGTCTCG
	281 TTACTGTGCGAGAGTGAATAACTGGAACGCCTTTGACCAGTGGGCCGTGGGCCGAAGTGCACTTTCTTCTGAGCTGACTC
	351 AGTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGGACAGCAGTCAGGATTACATGCCAAGGAGACACCCTCACAAG
	421 AGGACCCTGCTGTCTGTGCCTGGTACCAGCAGAAGCCAGGACAGGCCCCCTCCTGTCTTCTCTATGGTAAAGACAAG
	491 TTATTATGCGGGCCTGGTATCCCAGAGCCGATTCTCTGGCTCCAAGAAATATTGCTTCCTGACCATCACTG
	561 CGGCCCTCAGGGCCGGAGGATCCCGGGGCTGACTTTTACTGTGAGTTCCCGGGTACCGTTGGGTACCGTTTGTCTT
	631 GGGGCTCAGGGCTGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
	701 CGGGGCTGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv20 56	1 GAAGAAGCCTGGGTCCTCGGTGAAGGCTTCTGGAGGCACCTTCACCAGCTATGCAATC
	71 AGTTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGAGGGTTCATCCCTGTATTTGGCACAG
	141 CAAACTACGCCACAGAAGTTGCAGGGCAGAGTCACGATCACCGCGGACGAATTCCATGACCACAGTGTACAT
	211 GGAGCTGAGTAGCCTGACCTCTGAAGACACGGGGCGGTGTATTACTGTGCGAGGAATCTGTGCGAGGACGTGGGCCTGGCC
	281 CGTCGCGATGAATAACTACTATTACTACGATGGACGTCTGGGGCCAAGGGACAATGGTCACCGTCTCGAGTG

	351	GAGGCGGCCGGTTCAGGCGGGGAGGTGGCTCTCGGCGGTGGCGAAGTGCACAGTCTGTGCTGACTCAGCCACC
	421	CGCAGCGTCTGGGACCTACGGGCCAGAGAGATCCCAGGAGCGGCCCCCCACGTCGTCATCTATAATAATGATCAGC
	491	AATTATGTTTACTGGTACCGGTCCCTGACCGATCTCTGGCTGCTCCCGCTCCTCGGCCCATCAGTGG
	561	GCCCCTCAGGGGTCCGAGGATGAGGCTGATTATTGTTCCACATGGGATGACACCCTGAGTGGTTATATCTTC
	631	GCTCCGGTCCGAGGATGAGGCTGATTATTGTTCCACATGGGATGACACCCTGAGTGGTTATATCTTC
	701	GGAGTTGGGACCAAGGTCACCGTCCTAGGTGCGGCCGC
scFv21	57	
	1	CAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGATTCACCTTTAGCAGCTATGCCATGAGCT
	71	GGGTTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGTTATTAGTGGTGGTAGCACATA
	141	CTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATCTGCAA
	211	ATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCGAGAGATTGGAGATGGCAACAATTTG
	281	GGGGCTGGTTCGCACCCCTGGGCCGGGAACCCTGGTCACCGTCTCGAGTGGAGGCGGTTCAGGCGG
	351	AGGTGGCTCTGGCGGTGGCGGATCACATGCCAAGGAGTGCACATGCCAAGCCTTTAGCAGCTATGCCATGAGCT
	421	TTGGACAGACAGTCAGGATCACATGCCAAGGAGCAACCTCAGAAGTTTTTCTGACTCTGGTACCAGC
	491	TGAAGCCAGGACAGGCCCCTGTACTTGTCATCTATGGTAAGAACAACCGGCCCTCAGGGATCCCAGACCG
	561	ATTCTCTGCCTCCCAGCTCAGGAAACACAGCTTCCTTGGCCATCACTGGGCTCCAGGCTGAAGATGAGGCT
	631	GACTACTACTGTGAACTCTGGGGACCAGTCAGTGCTTATGTGTTCTTCGGAACTGGGACCAAGGTCACCG
	701	TCCTAGGTGCGGCCGC
scFv22	58	
	1	GGTCTTCGGTGAAGGTCTCCTGCAAAATTTCCGGAGGCAATCTCAATAGGCTTACTGTCACCTGGGTGCG
	71	ACAGGCCCCTGGACAAGGCCTTGAGTGGGTGGGCAGGATTCTTCCCGACTCAGTAAATCAGTCGTGAAG
	141	TTCCAGCGCAGACTCAAACTGACCTCTGACACTTCCACGCGCTATTTAGAACTGAGGAGCCTGA
	211	AATCTGAAGACACGGCCGTCTATTATTGTGCGGCGTCATCTAAAATAGGCTTCCAGTTGGGGAGCTCGA
	281	CTACTGGGGCCGGGGCCACCGTCTCGAGTGGAGGCGGCCGCGTCTGCTCACCCCGGCGGAGGGTCA
	351	GGTGGCGGAAGTGCACAGTGCACAGTGTCGTGACGCAGCCGCCCTCAGCCTACCCCGGGCAGAGGGTCA
	421	CCATCTCTTGTTCTGGAAGCAGTTCCAACATCGGAAGAAATTATGTCTACTGGTACCAGCAGCTCCCAGG
	491	AACGGCCCCCCAACTCCTCGTCTATAACAATAATCAGCGGCCCTCAGGGGTCCCGGTTCCGGCTCTGGC
	561	TCCAAGTCTGGCACCTCAGCCTCCCTGGCCATCAGTGGGCTCCAGGCTGAGGATGAGGCTGATTATTACT
	631	GTTCAACATGGGATGACAGCCTGAGTGGTGTATTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGC
	701	GGCCGC
scFv23	59	
	1	CACCTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTCTCAGCT
	71	ATTAGTAGTAGTGGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACA
	141	ATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGC
	211	GAGAGGTAGACGTCTGGGGGCCGGGGCACCGGGGAGTTAGACCCGGGCGGAGGCGGGAGGTGGCT
	281	ATGGACGTCTGGGGCCGGGGCACCGTCACCTTTCCATGTGCTCAGCGGAGGCGGTTCAGGCGGAGGTGGCT
	351	CTGGCGGTGGCGGAAGTGCACTTTCGTGCTGACTCAGCCCTCAGCGCTGTCTGCTCCCTCAGCTCCAGGACAGA
	421	TAGGGTCACCATCTCTTGTTCTGGAAGCAGCTCCAACATCGGAAGTAATTATGTATACTGGTACCAGCAG
	491	CTCCCAGGAACGGCCCCCAAACTCCTCATCTATAGGAATAATCAGCGGCCCTCAGGGGTCCCTGACCGAT

	561	TCTCTGGCTCCCAAGTCTCGGCCACCTCAGCCTCCCTGGCCATCAGTGGCTCCGGTCCGGAGGATGAGGCTGA
	631	TTATTACTGTGCAGCATGGGATGACACCCCTAAGTGGTGTCCTATTCGGCGGAGGACCAAGCTGACCGTC
	701	CTAGGTGCGGCCGC
scFv24	60	
	1	GCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGCTATTAGTGGTAG
	71	TGGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAAGAAC
	141	ACGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGCGAGAAATACAG
	211	GAAAGGGCATTACTTTGGTTCGGGGAGTATATTGTCAGGACTGTGACCGCAGTTCTACATCCCGCATGGA
	281	CGTCTGGGGCCAGGGCACCCTGGTCACCGTCTCGAGTGAGGCGGTTCAGGCGGAGGTGGCTCTGGC
	351	GGTGGCGGAAGTGCACAGGCTGTGCTGACTCAGCCGTCTCGACTCAGCGAAGGAATTATGTAGATTGGTCA
	421	CCATCTCTTGTTCTGGAAGCACTCCAACATCGGAAGTAATAATAAGCGGCCCTCAGGGGTCCCAGCTCCCAGG
	491	AACGGCCCCCAAACTCCTCATCTATAGGAATAATAAGCGGCCCTCAGGGGCTCCGGTCCGGAGGATGAGCTGGC
	561	TCCAAGTCTGGCCACCTCAGCCTCCCTGGCCATCAGTGGCTCCGGTATTCGGCGGAGGACCAAGCTGACCGTCCTAGGTGC
	631	GTGCAGCTTGGGATGACAGCCTGAGTGGTTGGGTATTCGGCGGAGGACCAAGCTGACCGTCCTAGGTGC
	701	GGCCGC
scFv25	61	
	1	AGGCTTGGTACAGCCTGGGGGGTCCCCGAGACTCTCCTGTGCAGCCTCTGGATTCACCTTTAGCAGCTAT
	71	GCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGCTATTAGTGGTAGTGGTG
	141	GTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAAGAACACGCT
	211	GTATCTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGCAAAAGATATGGGATAC
	281	AGTTATGGTTACGGGACGAGGGGCCTCTTTGACTACTGGGGCCAGGGAAGTGCACAGTCTCGTCACCGTCTCGAGTG
	351	GAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGGGCGGAGGTGGCACCGTCTGTCGTGACGCAGCCGCC
	421	CTCAGCGTCTGGGGCCCCCGGGGACAGAGATCACCATCTCTTGTTCCGGAAGCACCTTCAACATCGGGAGA
	491	AATTATGTTGACTGGTATAAACAACTCCCCGGAACGCCCCCCAAACTCTTCATCTATAAGAATGATCAGC
	561	GACCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGCCACTTGGGATGACAGCCTGGTCGTAAGTGG
	631	ACTCCGCTCCGAGGATGAGGCTGATTATTACTGTGCGGGGATGACAGCCTGAGTGGTGTCCGGTGTTC
	701	GGCGGGGGACCAAGGTCACCGTCCTAGGTGCGCGGCCGC
scFv26	62	
	1	TGGAGTCTGGGGGAGGCTTGGTACAGCCTGGGGGGTCCCTGTGCAGACTCTCCTGTGCAGCCTCTGGATTCAC
	71	CTTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCACGGGAAGGGGCTGGAGTGGGTCTCAGCTATT
	141	AGTGGTAGTGGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATT
	211	CCAAGAACACGCTGTATCTGCAAATGAACAATTTGGGGGCTGGTTCGGGAGGGCCGTGTATTACTGTGCGAG
	281	AGATTGGAGATGGCAACGGCGGGGTCCGGTGGCTCTGGGGGACCCCCCTGGGGCCGGGAGTCGACACTCACGC
	351	AGTGGAGGCGGGGTTCAGGCGGAGGTGGCTCTCCGGGGGACAGAGAGCCACCCTCTCCTGCAGGGCCAGTCTCACGC
	421	AGTCTCCAGCCACCCTGTCTGTTGGTACCAGCAGAAACCTGGCCAGGCTCCCATCTTTGGTGCATCC
	491	TGGTGGCAACTTAGCCTGGTACCCAGCCAGTTCAGTGGCAGTGGATCTGGGACAGACTTCACTCTCACCATCA
	561	ACTAGGGCCTCTGGGGTGTACCCCAGCCAGGTTCAGTGGCAGTGGATCTGGGACAGAGTTCACTCTCACCATCA

Fig. 11

scFv27	631	GCAGCCTGCAGTCTGAAGATTTGCAGTTTATTACTGTCAGCAGTATAATAACTGGCCTCCATGGACTTT
	701	CGGCCAAGGGACACGACTGGAGATTAAACGTGCGGCCGC
scFv28	1	ACAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGATTCACCTTTAGCAGCTATGCCATGAGC
	71	TGGGTCCGCCAGGCTCCAGGGAAGGGCCTGGAGTGGGTCTCAGCTATTAGTGGTGGTAGCACAT
	141	ACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATCTGCA
	211	AATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGCGGGGCGAGGGACCACGTCACCGTCTCGAGTGGCTGGA
	281	ACTACGTACTACTACTACGGTATGGACGTCTGGGGCCGGTGGACCGTGGTCGCCAGCCCGTCCGCCCTCAGC
	351	GCGGTTCAGGCGAGTGCTCTGGCGGTGGCGGAAGTGCACAGTCTGTGCTGACTCAGCCGCCCTCAGC
	421	GTCTGGGGCCCCCGGGCAGAGGATCACCATCTCTTGTTCCGGAAGCACCTTCAACATCGGGAGAAATTAT
	491	GTTGACTGGTATAAACAACTCCCCGGAACGGCCCCCAAACTCTTCATCTATAAGAATGATCAGCGACCCT
	561	CAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCCCTGGTCTGGTAAGTGGACTCCG
	631	CTCCGAGGATGAGGCTGATTATTACTGTCTGACTTGGGATGACAGCACCTGAGTGGTCCGGTGTTCGGCGGG
	701	GGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv29	1	GGCCTCTGGATTCGGCCTCAATGGCTATGAAATGCATTGGGTGCGCCAGGCCCCCGGACAAAGGCTTGAG
	71	TGGCTGGGCCGGATCAACGCTGCCATTGGCGACACGCTATTCAAGGGAGTTCCAGGATAGAGTCTCCA
	141	TTACCAGAGACATGTCCGGAACACAGTCTACATGGAGATGAGCAGTGAGATTGAAGACACGGCTGT
	211	TTATTATTGTGTGAGATTCCACGATTCCACGATTGTAATAGTGCCACCTGTCAGCCCCCTTTTGACCAC
	281	TGGGGCCAAGGGAACCCTGGTCACCGTCTCGAGTGGAGGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTG
	351	GCGGAAGTGCACTTTCTTCTGCAGCTCAGACCCTGTCTGCTGGCTCCTTGGGACACAGTCAG
	421	GATCACATGCCAAGGAGACAGCGTCAGACTATTCTGACAGTGGTACCGGCAGAAGCCAGGGCAGGCC
	491	CCTGTTATTGTCATGTATGATGTAACACCGCCGGCTCAGGGATCCCAGGCGTCGGGAGAGATGAGGCTGACTATTATTGTAACTC
	561	CAGGAAACACAGCTTCCTTGACCATCTGGTCTAACCATCTGGTCTTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
	631	CCGAGACAGTAGTGTGACTTCTTCGGCGGAGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv30	1	GTACAGCCTGGGGGGGTCCCTGAGACTCTCTGTGCAGACTCTCTGGATTCACCTTTAGCAGCTATGCCATGA
	71	GCTGGGTCCGCCAGGCTCCGTGAAGGGCGGTTCACCATCTCCAGAGACAATTCCAGAGAGTGGTGGCAGCAC
	141	ATACTACGCAGACTCCGTGAGAGCCGAGGACACGGCCGTTCCAGAGAATTCCAAGAACACGCTGTATCTG
	211	CAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGGGGCGTGGGGACTACTGCGGCGTCGGGGACGCG
	281	GAGGTGGGAGCTACTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCAATGGTCACCGTCTCGAGTGG
	351	AGGCGGCGGTTCAGGGGCCCCCGGGGACCCCGGGCTGGCTCTGGCGGTGGCCACATCGGGAAGGA
	421	TCAGCGTCTGGGGCTCTGGGGACCCCGGGCAGCAGGGTCCCCACCATCTCTTGTTCTGGAAGCAGTTCCAACATCGGAAGGA
	491	ATTATGTATACTGGTACCAGCAGCTCCCAGGAACGGCCCCCAAACTACTCATCTATAGAAATAATCTGCG
	561	GCCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCCCTGGCCATCAGTGGG
	631	CTCCGGTCCGGAGGATGAGGCTGATTATTACTGTGCAGCAGTTGCACCCTGAGTGGTGTGATTCG
	701	GCGGAGGGACCAAAGCTGACCGTCCTAGGTGCGGCCGC
scFv30	1	CGGAGGTGAGGAAGCCTGGGGCCTCAGTGGAGGATTCCTGCAAGGCTTCTGGATTCACGTTCACTAGTTA
	71	TCTATTCCATTGGGTGCGCCAGGCCCCCGGACAAAGGCTTGAGTGGGATGGGGTGGATCAACGCTGGCAAT
	141	GGAAACACAAAATATTCACCGGAGTTACCCTTACCGGGACACATCCACGAGCACAA

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     211  CCTACACATGGAGCTGAGCAGCCTGACATCTGAGGACACGGCTGTTTATTACTGTGCGAGAGATCAGGTGTT
     281  CTATGAGAGTGGTTCTTACTACATACGCCCTTCTTTGACTTCTGGGGCCGCAGGGGACCCTGGTCACCGTC
     351  TCTTCAGGTGGAGGCGGTTCAGGCGGAGGTGGCTCAGGCGGTGGCGGATCGGACATCCAGATGACCCAGT
     421  CTCCTTCCACCCTGTCTGCATCTATTGGAGACAGAGTCACCATCACCTGCCGGCCAGTGAGGGTATTTA
     491  TCACTGGTTGGCCTGGTATCAGCAGAAGCCAGGGAAAGCCCCTAAACTCCTGATCTATAAGGCCTCTAGT
     561  TTAGCCAGTGGGGCCCCATCAAGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCA
     631  GCCTGCAGCCTGATGATTTTGCAACTTATTACTGCCAACAATATAGTAATTATCCGCTCACTTTCGGCGG
     701  AGGGACCAAGCTGGAGATCAAACGTGCGGCGCGC
```

scFv31 — 67

```
       1  TGGTACGGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGCCTCTCTGGATTCACCTTTGATGATTATGGCAT
      71  GAGCTGGGTCCGCCAAGCTCCAGGGAAGGGGCTGGAGTGGGTCTCGTATTAATTGGAATGGTGGTAGC
     141  ACAGGTTATGCAGACTCTGTGAAGGGCCGATTCACCATCTCCAGAGACAACGCCAAGAACTCCCTGTATC
     211  TGCAAATAAACAGTCTGAGAGCCGAGGACACGCCGTGTATTACTGTGCAAGAGGCGGTATGCGTTGGA
     281  TTATTGGGGCAGAGGGACCAATGGTCACCGTCTCGAGCTGACTCAGGACCCTGCTGTGTCTGTGGCCTGGC
     351  GGTGGCGGAAGTGCACTTTCTTCTGAGCTGACTCAGGACCCTGCTGTGTCTGTGGCCTGGG
     421  CAGTCAGGACATAACTTGTCAGGGACACCTCGACACAAATATTATGCAAACTGGTATCAACAGAAGCCTGG
     491  ACAGGCCCCTCTACTTGTCTTCTTCTTTTCTGACCATCACTGGGCCCTCAGGGCTCCAGGCGTTTCTCTGGC
     561  TCCAACTGCGGAACACAGCTTCCTGACCATCAATCCATCCTAATCGGTACTAATCGAATCGTTCCTAGGTGC
     631  GCAACTCCCGGGAAATCGGTACTAATCGAATCGTTCCTAGGTGC
     701  GGCCGC
```

scFv32 — 68

```
       1  TTGGTTCAGCCTGGAGGGTCCCTGAGACTCTCCTGTGCAGCCGCTGGATTCACCTTCAGTACTTTTGAAA
      71  TGAATTGGGTCCGCCAGGCCCCAGGGAAGGGGCTGGAGTGGGTTTCATATATTAGTGGTAGTGGTCATGC
     141  CATATACTACGCAGAGTCTGACCTGAGAGGCCGATTCACCATCTCCAGAGACAACGCCAACAACTCACTGTAT
     211  CTGCAAATGAACAGTCTGAGAGCCGAGGACACGGCTGTTTATTACTGTGCGAGAAAAGTACCAGCTAC
     281  TACTTGGCAAGTACGACTACACGGTATGGACGTCTGGGGCCGGGCGAAGTGCACCGTCTCGAGTGGAGG
     351  CGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGGAAGTGCACTGCCTCGTGCTGACTCAGCCACCCTCA
     421  GCGTCTGGGACCCCCGGGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAGCTCCAACATCGGAAGTAATA
     491  CTTTAAACTGGTACCAGCAGCTCCCAGGAACGGCCCCCAAACTCCTCATCTATAGTAATAATCGGCTGGCC
     561  CTCAGGGGTCCCTGACCGGTTCTCTGGCTCCAAGTCTGGCACCTCCGCCATCAGTGGGCTC
     631  CAGTCTGAGGATGAGGCTGATTACTACTGTGCAGCATGGGATGACAGCAGCCTGAATGGCTGGGTGTTCGGCG
     701  GAGGGACCAAGGTCACCGTCCTAGGTGCGCGGCGC
```

scFv33 — 69

```
       1  AGGGCCCTCTGGGGGACCTCTGGGGTGCTTTCAGCTGGTGCGACAGGCCCCTGGACAGGGGCTTC
      71  AGTGGATGGGAGGGATCATCCCTCTCTTTTGGTGCAGCAAATACCGCACAAAAGGTCCGGGCCGACTCAC
     141  GATTACCGCGGACGAGTCCACGGGACGTCTTACAGGGAGCAGCCTGAGACAAGCTGACACGGCC
     211  GTTTATTTCTGTGCGACTAACGGACAGAGCGAGGTCGCCACCCGGCTACTACTACGGGCATGGACGTCTGGG
```


Fig. 11

	281	GCCGAGGCACCCTGGTCACCGTCTCGAGTGGAGGCGGCGGTTCAGGCGGCGGAGGTGGCTCTGGCGGTGGCGG
	351	AAGTGCACAGTCTGTGTTGACGCAGCTGCCCTCGGGGCTCCAGCGTCGGGGCCCCGGGCAGAGGATCACCATCTCT
	421	TGTTCCGGAAGCACCTTCAACATCGGAGAAATTATGTTGACTGGTATAAACAACTCCCGAACGGCCC
	491	CCAAACTCTTCATCTATAAGAATGATCAGCGACCCTCAGGGGTCCCTGCCCGATTCTCTGGCTCCAAGTC
	561	TGGCACCTCAGCCTGACCCTCCCTGGTCATCGTCGTCGTAAGTGGACTGGACTGAGGCTGAGGCTGATTATTACTGTGCTGACT
	631	TGGGATGACAGCCTGAGTGGTGTCCGGTGTTCGGCGGGGGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv34		
	1	GCCTGCAAGGTTTTGGTTACACCTTCGTCGATCATGGAATTAGTTGGGTGCGACAGGCCCCTGGACAAG
	71	GGCTTGAGTGGATGGGATGGATCAACACTCACGACGTCACCAAACTATGCACAAAGACACAGGCCAG
	141	ACTCACCATGACCACCAGAGTGCCTCCCATTAATACTTCCTACATGGAGCTGCCGAGCCTGACATCTGACGAC
	211	ACGGCCGTCTATTATTGTGCCCGGGGGGAGAGACTCGGACGTCGGGGCCAAGAACCCTGGTCACCGTCTCGAG
	281	ACAATGGATATCCCTATTACTCCTATTACTCCTGGTCTGGACGTGGCGGTCTGGCGGAGGTGGCTCTGGCGGTGGCGG
	351	TGGAGGCGGCGGTTCAGGCGGACCCCCGGGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAGCTCCAACATCGGAA
	421	TCCTCAGCGTCTGGGACCCCGGGCAGCAGTCCCAGGAACGGCCCCCAAACTCCTCATCTATAGGAATAATCA
	491	GTAATTATGTATACTGGTACCAGCAGCTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCTGGGCCATCAGT
	561	GCGGCCCTCAGGGGTCCCTGACCGATTCTGATTATTACTGTGCAGCATGGGATGACAGCCTGAGTGGTTGGGTGT
	631	GGGCTTCCGGTCCGAGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
	701	TCGGCGGAGGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
scFv35		
	1	AGCCTGGGGCCCTCAGTGAAGGTTTCCTGCAAGGCATCTGGATACACCTTCACCAGCTACTATATGCACTG
	71	GGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACCCTAGTGGTGGTAGCACAAGC
	141	TACGGACAGAAGTTCCAGGGCAGAGTCACCATGACCAGGGACACGTCCACGAGACACAGTCTACATGGAGC
	211	TGAGCAGCCTGAGATCTGAGGACACGGCCGTGTATTACTGTGGGCGGCCAGAGGTTCGGGCGCCAGAATGGTTCG
	281	GGGAGTTATTATTAGACCCCTACGGTATGGACGTCTCGGGGCCGAAGTGACCCTGGTCACCGTCTCGAGTGGA
	351	GGCGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGAAGTGCACAGTCTGTGCTGACTCAGCCACCCTCT
	421	CAGCGTCTGGGACCCCCGGGCAGAGGGTCACCATCTCTTGTTCTGGAAGCAGCTCCAACGTCGGAAGTAA
	491	TTATGTATCCTGGTATCAGCAGTTCCCAGGAACGGCCCCCAAACTCCTCATCTATAGGAATAATCAGCGG
	561	CCCTCAGGGGTCCCTGACCGGTTCTCTGGCTCCAAGTCTGGCATTTCAGCCTGGCCATCAGTGGGC
	631	TCCGGTCCGAGGATGAGGCTGATTTTACTGTGTAGCAGCATGGGATGACAGCCTGAGTGGTTGGGTGT
	701	AACTGGGACCAAGGTCACCGTCCTAGGTGCGGCCGC
scFv36		
	1	GGAGTCTGGGGGAGGCGTTGGTACAGCCTGGGGGTCCCTGAGACTCTCTCCTGTGCAGCCTCTGGATTCACC
	71	TTTAGCAGCTATGCCATGAGCTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCTCAGCTATTA
	141	GTGGTAGTGGTAGCACATACTACGCAGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTC
	211	CAAGAACACGCTAGGGGTGTATCTGCAAATGAACAGCCTGAGATATGGGCGTGTATTACTGTGCGAAA
	281	GGTGGGGACTAGGGTGACCACCGGGGGTGGTTTGATATATGGGCCGGTCTGGCGGTGGCTGCTGACTCGA
	351	GTGGAGGCGGCGGTCTCGGGGCCCCGGGCAGAGGATCACCATCTCCGGAAGCACCTTCAACATCGGG
	421	CCCCTCAGCGTCTGGTATAAACAACTCCCCGGAACGGCCCCCAAACTCTTCATCTATAAGAATGATC
	491	AGAAATTATGTTGACTGGTATAAACAACTCCCGAACTCTTCATCTATAAGAATGATC
	561	AGCGACCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCCCTGGTCGTAAG

Fig. 11

631 TGGACTCCGCTCCGAGGATGAGGCTGATTATTACTGTCTGACTTGGGATGACAGCCTGAGTGGTGGTCCGGTG
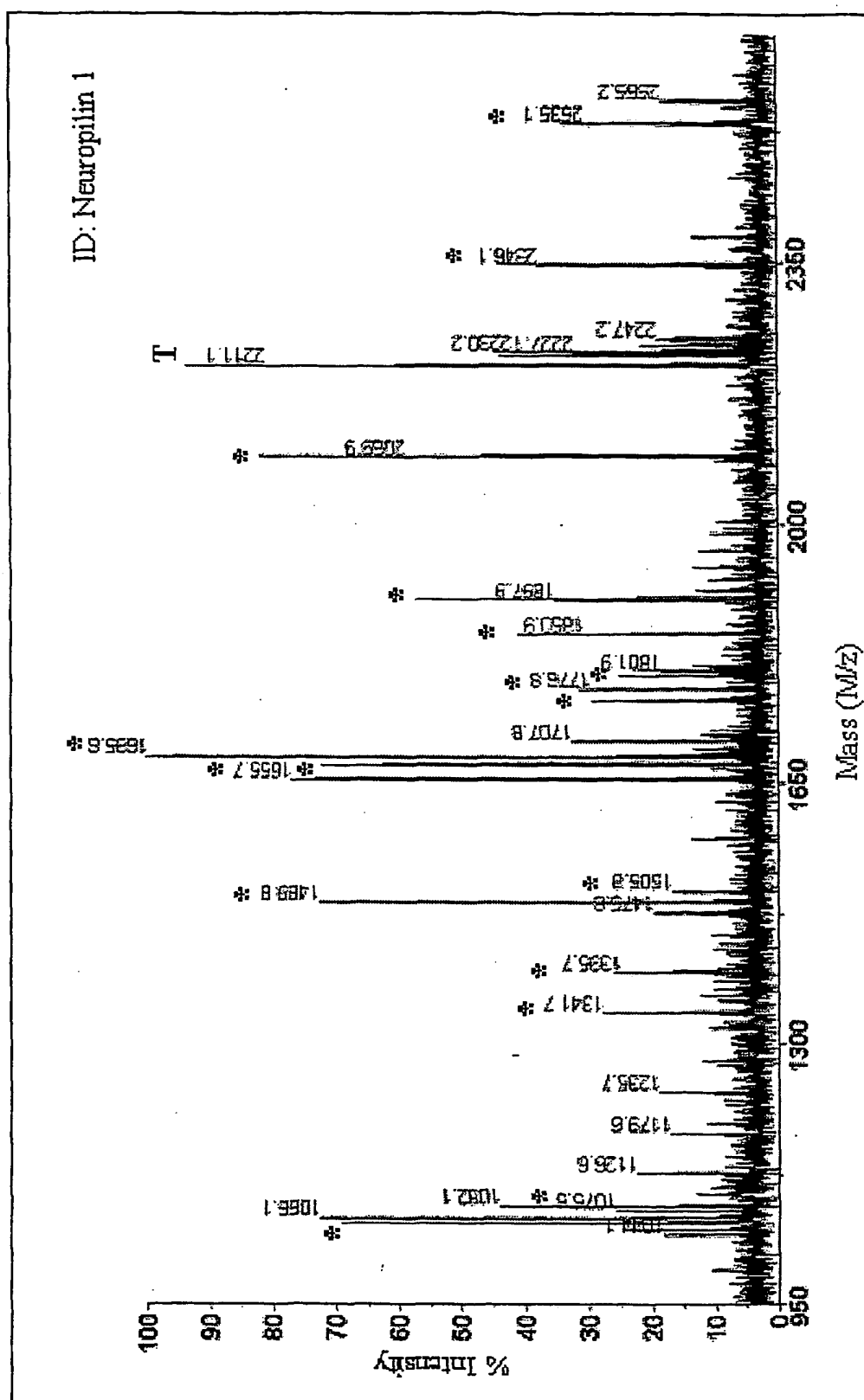701 TTCGGCGGGGGGACCAAGCTGACCGTCCTAGGTGCGGCCGC
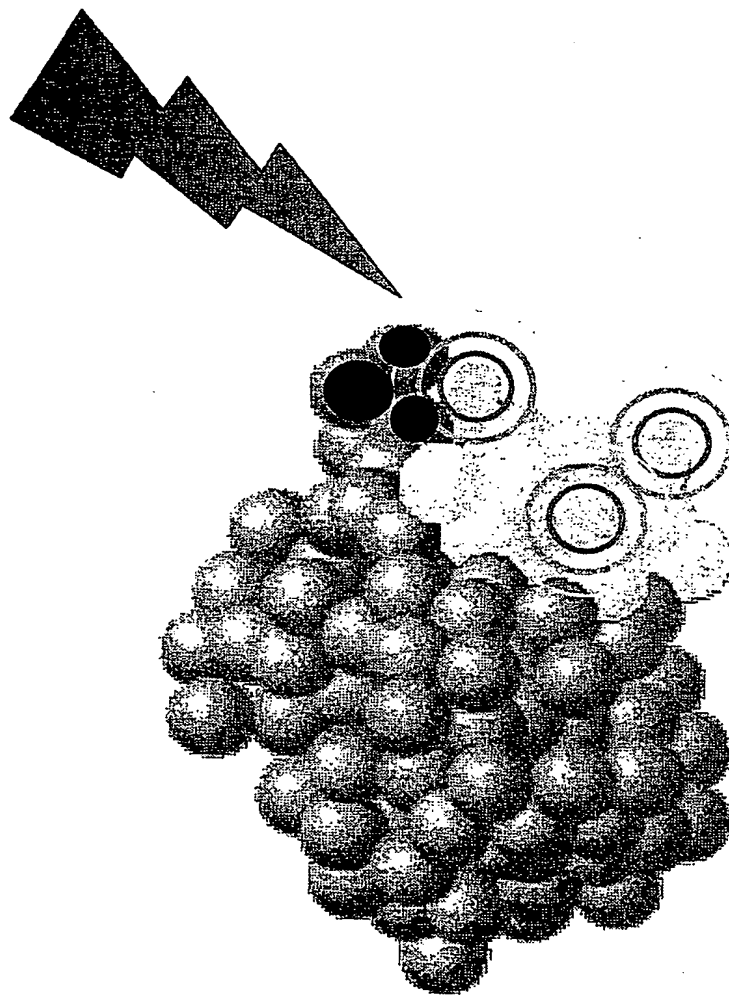
Fig. 11

Fig. 12

Fig. 13

Fig. 14c

Fig. 14b

Fig. 14a

10/539402

Tab. 1: Inhibition of tube formation

scFv	Average inhibitory effect
control (10% PBS)	0.8
scFv4	1.7
scFv5	1.6
scFv7	2.4
scFv8	2.4
scFv9	1.5
scFv11	2.2
scFv12	1.9
scFv13	2.1
scFv15	1.9
scFv16	1.6
scFv18	2.0
scFv19	1.6
scFv20	1.3
scFv21	2.0
scFv22	1.5
scFv23	2.1
scFv24	1.7
scFv25	2.3
scFv26	2.1
scFv27	2.3
scFv28	2.2
scFv29	2.1
scFv30	1.8
scFv31	2.4
scFv32	1.7
scFv33	2.1
scFv34	1.5
scFv35	1.4
scFv36	1.9
scFv7*	2.3
scFv8*	2.5
scFv13*	2.5
scFv25*	2.6
scFv26*	2.4
scFv28*	2.1
scFv31*	2.3
scFv33*	2.1
anti-alpha-2-integrin	2.3
anti-NP-1	1.1

Fig. 15
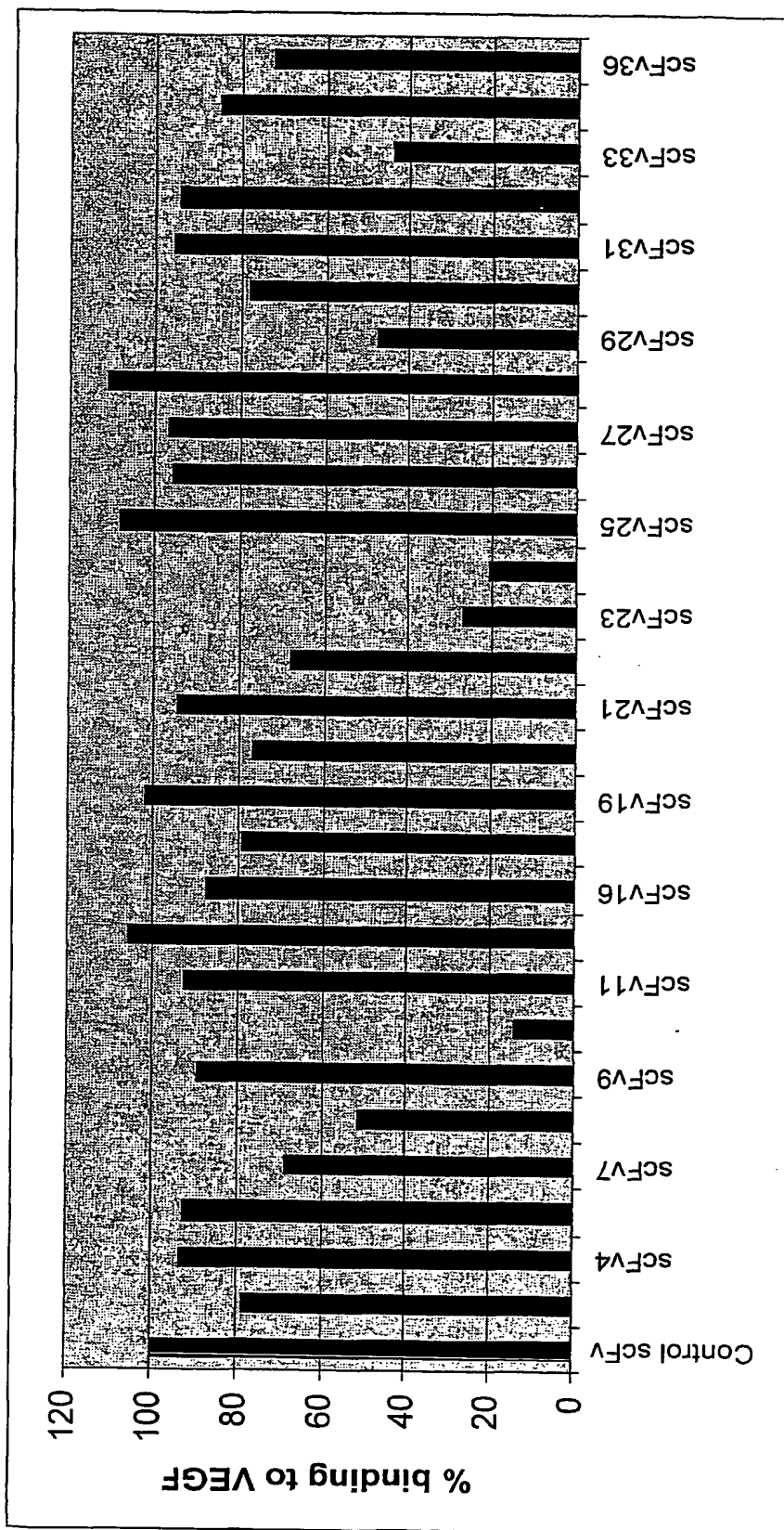
Fig. 16
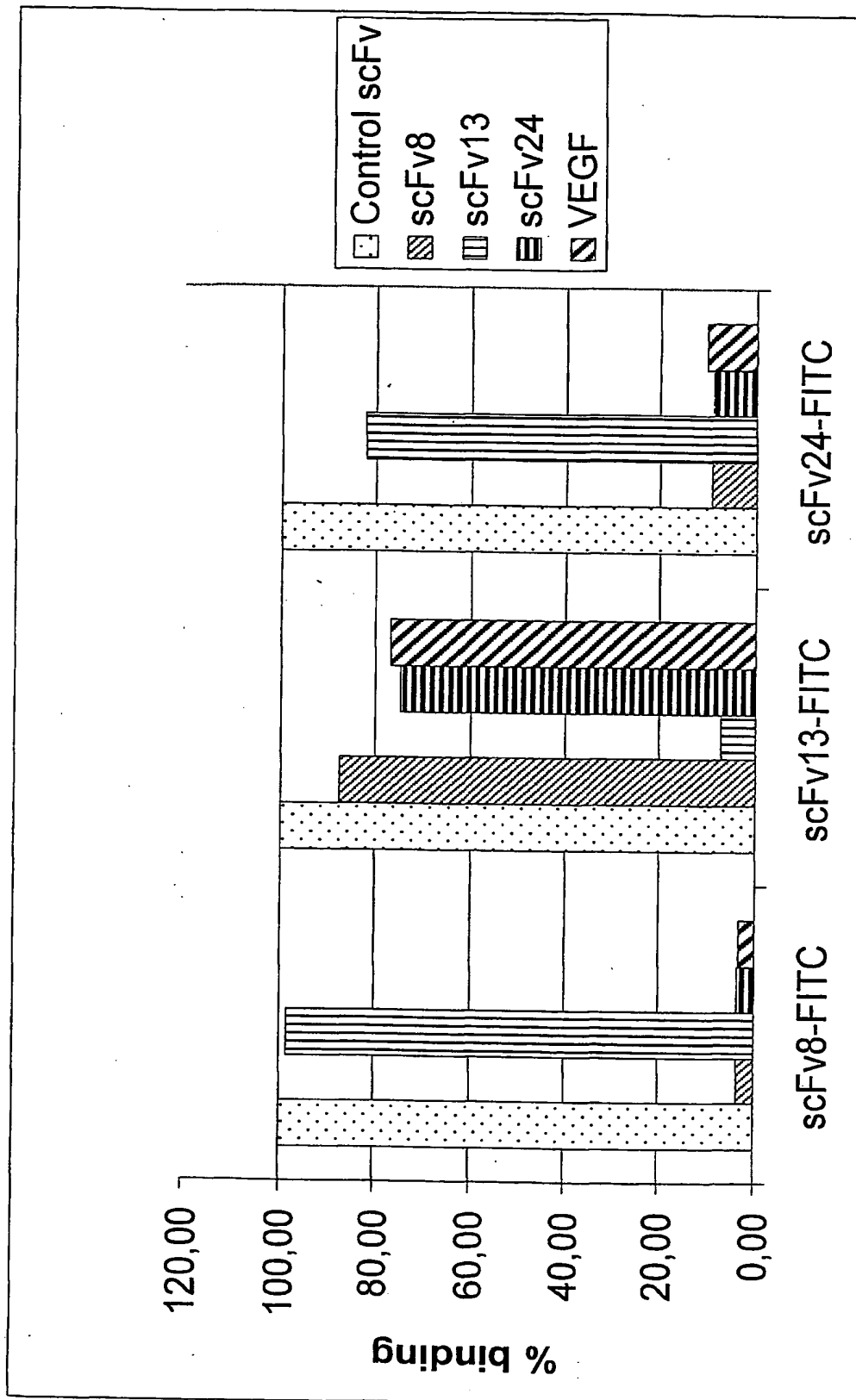
Fig. 17

Tab.2   Transendothelial Invasion assay

scFv	Inhibition of Invasion
scFv26	+
scFv27	+
scFv34	+
ScFv35	+

Fig. 18

Fig. 19